Sample handling in surface sensitive chemical and biological sensing: A practical review of basic fluidics and analyte transport

Norbert Orgovana,b, Daniel Patko b,c, Csaba Hos d, Sándor Kurunczi b, Bálint Szabó a,
Jeremy J. Ramsden e,f, Robert Horvath b,k

a Department of Biological Physics, Eötvös University, Pázmány P. sétány 1A, H-1117 Budapest, Hungary
b Nanobiosensorics Laboratory, Institute for Technical Physics and Materials Science, Research Centre for Natural Sciences (MTA TTK MFA), H-1121, Konkoly-Thege Miklós út 29-33, Budapest, Hungary
c Doctoral School of Molecular- and Nanotechnologies, Faculty of Information Technology, University of Pannonia, H-8200 Egyetem u. 10, Veszprém, Hungary
d Department of Hydraulic Machines, Budapest University of Technology and Economics (BME), 1521 Budapest, Pf. 91, Hungary
e Clore Laboratory, University of Buckingham, MK18 1EG, UK
f Centre for Molecular Recognition, Collegium Basilea (Institute of Advanced Study), Hochstrasse 51, CH-4053 Basel, Switzerland

ARTICLE INFO
Available online 13 April 2014

Keywords:
Fluidic systems
Analyte transport
Dead volumes
Label-free detection
Optical biosensors
Fluid handling of live cells

ABSTRACT
This paper gives an overview of the advantages and associated caveats of the most common sample handling methods in surface-sensitive chemical and biological sensing. We summarize the basic theoretical and practical considerations one faces when designing and assembling the fluidic part of the sensor devices. The influence of analyte size, the use of closed and flow-through cuvettes, the importance of flow rate, tubing length and diameter, bubble traps, pressure-driven pumping, cuvette dead volumes, and sample injection systems are all discussed. Typical application areas of particular arrangements are also highlighted, such as the monitoring of cellular adhesion, biomolecule adsorption–desorption and ligand–receptor affinity binding. Our work is a practical review in the sense that for every sample handling arrangement considered we present our own experimental data and critically review our experience with the given arrangement. In the experimental part we focus on sample handling in optical waveguide lightmode spectroscopy (OWLS) measurements, but the present study is equally applicable for other biosensing technologies in which an analyte in solution is captured at a surface and its presence is monitored. Explicit attention is given to features that are expected to play an increasingly decisive role in determining the reliability of (bio)chemical sensing measurements, such as analyte transport to the sensor surface; the distorting influence of dead volumes in the fluidic system; and the appropriate sample handling of cell suspensions (e.g. their quasi-simultaneous deposition). At the appropriate places, biological aspects closely related to fluidics (e.g. cellular mechanotransduction, competitive adsorption, blood flow in veins) are also discussed, particularly with regard to their models used in biosensing.

© 2014 Elsevier B.V. All rights reserved.

Contents
1. Introduction ............................................................... 2
2. Short overview of the basic theory of fluidics and analyte transport ............................................................... 3
2.1. Fluid handling in cuvettes and cylindrical connecting tubes — flow rates ......................................................... 3
2.2. Diffusion of analytes: the diffusion boundary layer ............................................................... 4
2.3. Transport to, adsorption on, and desorption from the sensor surface ............................................................... 4
2.3.1. Adsorption model for single-component solutions ............................................................... 4
2.3.2. Desorption from the sensor surface — readsoption and rebinding ............................................................... 5
2.3.3. Adsorption from complex, multicomponent analyte solutions ............................................................... 5
2.4. Fluid handling in cell-based assays ............................................................... 5
2.4.1. Adhesion assays: general considerations, and cell deposition in an environment without continuous flow ............................................................... 5
2.4.2. Mechanotransduction assays: cells in a flow environment ............................................................... 6
2.4.3. Signaling assays: the challenge of substance addition to pre-attached cells ............................................................... 6
3. Experimental: materials and methods ............................................................... 7
3.1. Optical waveguide lightmode spectroscopy ............................................................... 7

* Corresponding author.
E-mail address: horvath@mfakfki.hu (R. Horvath).

http://dx.doi.org/10.1016/j.cis.2014.03.011
0001-8686/© 2014 Elsevier B.V. All rights reserved.
Table of notation

A Cross section of a tube
α Constant of proportionality between N and analyte concentration
C Constant characterizing the geometry of a cuvette
c_0 Bulk concentration
c_ν Analyte concentration in the vicinity of an adsorbing surface
Γ Surface mass density of reversibly adsorbed molecules
Γ′ Surface mass density of irreversibly adsorbed molecules
d Diameter of a tube
D Diffusion coefficient
δ Thickness of the diffusion boundary layer
F Volumetric flow rate
φ Available area function
g Gravitational constant
k_B Boltzmann constant
k_d First order rate coefficient of desorption
k_i Rate coefficient of irreversible adsorption
k_r Rate coefficient of reversible adsorption
L Length of tubing
N Effective refractive index
μ Dynamic viscosity
n_c Refractive index of the cover layer (medium)
ρ_1,ρ_2 Pressures at the two ends of a tube
q Distance between the in- and outlet apertures of a flow-through cuvette
r Radius of a diffusing object
R Inner radius of a tube
Re Reynolds number
ρ Density of a fluid
ρ_c Mean density of a mammalian cell
ρ_m Density of an aqueous medium
S Unit area
t Time
t_s Time required to saturate a biosensor signal
T Absolute temperature
τ_w Wall shear stress
v Mean velocity of a fluid
v_0 Velocity of the leading edge
v_term Terminal velocity of a cell during sedimentation
V Unit volume
V_f Total sample volume of a cuvette
V_c Volume of the convective zone of a cuvette
V_u Volume of the unflushed regions of a cuvette
x Distance
z Distance between the point of fluid sample inlet and point of sensing in a cuvette

1. Introduction

Nowadays, the field of chemical and biological sensing goes well beyond the application of novel technologies to detect the presence of a given analyte. The range of table-top biosensor instruments is wide and their use helps many laboratories worldwide to answer important scientific and technological questions in biomaterials and surface sciences, drug development, biophysics, (bio)nanotechnology and cell biology [1], as well as supramolecular chemistry [2].

Biosensing technologies enabling label-free detection at a solid-liquid interface include surface plasmon resonance (SPR) (and SPR imaging, SPRi) [3–5], quartz crystal microbalance (QCM) [6], dual polarization interferometry (DPI) [7,8], grating-coupled interferometry (GCI) [9–12], the high-throughput compatible resonant waveguide grating (RWG) biosensing [13–15], and optical waveguide lighmode spectroscopy (OWLS) [8,16–18]. OWLS instruments are currently among the most sensitive commercial label-free biosensors that enable the monitoring of various processes accompanied by refractive index changes up to 100–200 nm above a sensor surface. OWLS permits real-time, high-sensitivity (adsorbed mass coverages as low as 10 pg/mm² can be detected) label-free detection with a typical temporal resolution of 10–20 s, which makes it an ideal research tool in several fields, including the online monitoring of protein adsorption [19–22], bacterial [23,24] and mammalian cell adhesion [25–30], supported lipid-bilayer structure [31,32], polyelectrolyte multilayer build-up [33–35], nanoparticle surface adhesion and assembly [36,37], and receptor–ligand interactions [38,39].

An outstanding advantage of surface-sensitive label-free biosensing technologies is that they allow the so-called “real-time monitoring”, thus generating kinetic data. Kinetic modeling based on such valuable data enables the molecular or cellular process under investigation to be quantitatively characterized in detail. For example, adsorption and desorption rates at model surfaces, or kinetic rates in affinity binding can be determined [40,41], which contribute to a better understanding of the mechanism underlying the investigated surface process. Kinetic analysis of experimental data is furthermore the key to a detailed characterization of cell adhesion [14], including the separation of the governing cellular processes from less relevant ones, and the quantification of the differences arising from different cell types, substrata, media and other environmental factors [14,42–46]. However, kinetic data is often misinterpreted. A common reason is that the temporal evolution of the concentration distribution of the investigated objects over the sensing area is not known, or is uncontrolled; the extent of these effects depend on the fluidic system used.

In the majority of applications, biomolecules or cells are investigated in aqueous solutions mimicking their native environment. OWLS measurements are usually carried out by sealing a cuvette containing the medium over the sensing area. Generating flow to ensure a continuous supply of fresh analyte, changing solutions, executing washing steps, etc. all require some kind of fluid handling system.
A variety of fluidic systems can be built from the wide range of commercially available accessories in order to optimize sample handling for a particular application. The simplest systems use a plain well without flow (henceforth “closed cuvette”). However, this simplest configuration may yield less informative data as compared to measurements when the liquid sample is continuously flowed over the sensing area [17]. More sophisticated fluidic systems, therefore, consist of pumps, sections of tubes of different lengths and diameters, the junctions between them, and a flow-through cuvette (Fig. 1). Other components such as bubble traps can be integrated as well [47].

However, in several applications, especially when working with costly analytes, the continuous flow of fresh material is not feasible. In these applications, some kind of sample injection system has to be used, preferably in combination with a flow-through system. This facilitates sample handling automation: the sample is introduced by aspiration or injection and subsequently guided to the cuvette. This technique is called flow injection analysis (FIA), which has many valuable features for on-line biosensor measurements [48]. Although FIA can be used to modify the sample matrix or for preconcentration or derivatization, biosensing has predominantly exploited the precise and reproducible fluid handling capability of the injection valves used with FIA [49,50].

Importantly, the fluidic system is a critical part of any biosensor setup, as it is the most common source of experimental errors, which endanger the reliability of the recorded data and may lead to serious misinterpretations. This is especially true when elaborate manipulations with the samples have to be made or when the kinetics of the processes are deeply interpreted. The present work overviews the most important theoretical and practical considerations involved in the design of a fluidic system. We aim at facilitating finding the best arrangement for a given application; and emphasizing how to obtain reliable data, which can be then subjected to detailed analysis. In the experimental part we focus on sample handling in OWLS measurements, but this study is equally applicable to other biosensing technologies in which an analyte in solution is captured at a surface and its presence is monitored. At the appropriate places, key biological phenomena closely related to fluidics (e.g., cellular mechanotransduction, competitive adsorption, blood flow in veins) are also highlighted to indicate how in vivo conditions can be better approximated and understood. After a brief description, the reader is referred to papers specifically dealing with these problems.

2. Short overview of the basic theory of fluidics and analyte transport

2.1. Fluid handling in cuvettes and cylindrical connecting tubes — flow rates

The supply of analyte solution is usually pumped through tubes to reach a tube-like cuvette placed over the sensing area. The nature of the flow is determined by the value of the dimensionless Reynolds number \( \text{Re} \), defined as the ratio of inertial forces \( (\text{due to the flow}) \) and viscous forces \( (\text{due to the internal friction of the fluid}) \):

\[
\text{Re} = \frac{\rho v^2/d}{\mu} = \frac{\rho v d}{\mu},
\]

where \( \rho, \mu \) and \( v \) are the density, dynamic viscosity and mean velocity of the liquid, respectively, \( d \) is the diameter of the tube, \( F \) is the volumetric flow rate and \( A \) is the cross-section of the tube [51]. Below a critical value of \( \text{Re} \), viscous forces dominate and the flow is laminar, i.e. the velocity depends only on the perpendicular distance from the fore-axis of the tube. In this case the evolving streamlines are parallel to the axis, and the stream volume can be split into coaxial layers in which the flow rates are the same (Fig. 1). The flow is surely laminar in a tube when \( \text{Re} < 2300 \), and turbulent when \( \text{Re} > 4000 \). In typical biosensing experiments the flow rates guarantee that \( \text{Re} \) is well below the former threshold, therefore only laminar flow occurs [52]. With the present focus on instrumental miniaturization (e.g., the creation of labs-on-chips) internal tubing diameter can be as small as 100 \( \mu \)m or even less; \( \text{Re} \) is then of order unity [53]. An important exception to the preponderance of laminar flow is when an atypically high flow rate is used in order to effectively flush the dead volumes of the cuvette by generating turbulence (see Section 4.2.2.1 and Fig. 6).

A number of practical aspects set upper and lower limits to the actual flow rate with which an analyte solution can be pumped during a biosensing experiment. The chosen rate determines how much sample is needed for one measurement and whether the investigated process will be adsorption- or transport-limited (see Section 2.3); in the latter case the resulting kinetic data will not be representative of the true surface process (as may happen in Biacore experiments using the popular dextran-coated surfaces [54]).

Expense or limited availability of a sample pushes one to use the least possible amount in a measurement and, as an initial strategy, the amount of analyte, during which the surface ad-

![Fig. 1. Schematics (not to scale) showing an OWLS device with a basic flow-through fluidic system (bottom); and the bottom half of the parabolic flow profile inside the cuvette (top). Any kind of surface process is investigated, the reliability of the kinetic data is severely dependent on the appropriate fluidic design and on a reproducible fluid handling strategy. At the bottom: an inlet tube connects the OWLS flow-through cuvette (mounted on the goniometer) with the sample reservoir on the left. In between, a bubble trap is seen. The constant flow of analyte is engendered with a peristaltic pump, and the outlet tube leads the overflow to a waste container. Although this is considered to be a basic fluidic system, its constituting elements and the way the liquid samples are handled may severely influence the kinetics of the surface process that has to be quantified. At the analyte transport and shear stresses in the cuvette. In biosensors, the engendered flow is typically laminar and, thus, the flow profile is parabolic. Close to the walls of the tubing or that of the cuvette, convection is negligible and solutions are exchanged via diffusion in a layer called the diffusion boundary layer (arrow to the left). An analyte (e.g., a protein) may adsorb reversibly or irreversibly to the surface, or may desorb from there (with corresponding rate coefficients of \( k_a, k_d \) and \( k_e \)); changing causes in the local refractive index, which is immediately detected. The biosensor also enables the presence and activity of living cells seeded on the sensor surface to be monitored, although generally only the bottom portion of the cell can be probed (i.e. the sensing depth – shown as a reddish layer, and marked with a red arrow to the right – is only 100–200 nm). OWLS in combination with appropriately designed flow-through fluidic systems can be used to apply well-defined shear forces \( F_s \) to cells attached to the sensor surface, and monitor their response (cellular mechanotransduction).]
of the sample first appears in the recorded data, the cuvette has already been perfectly flushed through by the sample solution (e.g., the liquids have been perfectly exchanged). The lower the flow rate, the more carefully this assumption needs scrutiny (Section 2.3).

In this review we consider only pressure-generated flow. If it is laminar, the Hagen–Poiseuille law states that

\[ p_1 - p_2 = \frac{8 \mu L}{\pi R^4} T, \]  

(2)

where \( p_1 \) and \( p_2 \) are the pressures at the two ends of the tube, and \( R \) and \( L \) are the inner radius and length of the tube, respectively. Based on the formal similarity of Eq. (2) to Ohm’s law, the multiparameter factor multiplying \( F \) on the right side of Eq. (2) is usually referred to as the “resistance” of the tube. Note how this factor depends on \( R \) and \( L \). If this resistance is changed following modification of the setup, it is strongly recommended to check whether the pump is powerful enough to generate the same flow rate as earlier.

Fluid dynamics and flow in a tube have important biological aspects, especially when considering flow in blood vessels. Note, however, that pulsatile flow of (the non-Newtonian) blood in the compliant vessels is far too complex to be characterized by the classical Poiseuille fluid dynamics (describing steady flow in a rigid tube of circular cross-section) [56] briefly introduced above.

2.2. Diffusion of analytes: the diffusion boundary layer

Surface-sensitive, high-performance label-free techniques require the sample of interest to be brought into close contact with the surface of the sensor (the planar waveguide in OWLS, RWG, and GCC; or the gold surface in SPR). During analyte transport two fundamental processes have to be distinguished: convection (sometimes called advection) and diffusion. The liquids in which the biological or chemical objects under investigation are dissolved or suspended are usually introduced above the sensor surface by pumps, engendering convection [51]. At the solid–liquid interface, however, the tangential flux is always zero and, therefore, another transport process, diffusion, dominates close to the surface. The importance of diffusion relative to other transport phenomena (convection, sedimentation) is strongly dependent on the nature of the suspended analyte objects: their size, buoyancy, 3D shape and structure are all important factors.

An object having a diffusion coefficient \( D \) will diffuse to an expected distance of

\[ \sqrt{\langle x^2 \rangle} = \sqrt{\frac{D}{q} t} \]  

(3)

from its starting point during elapsed time \( t \). \( q \) is a numerical constant which depends on dimensionality: it is 2, 4 or 6 for 1, 2, or 3 dimensional diffusion, respectively. Assuming spherical symmetry for the analyte object, the Stokes–Einstein equation can be used to calculate \( D \):

\[ D = k_B T / 6 \pi \eta r \]  

(4)

where \( k_B \) denotes the Boltzmann constant, \( T \) the absolute temperature, and \( r \) the radius of the object; if it is small the effect of gravity is negligible.

Diffusion plays an important role in other parts of the fluidic system as well as above the sensor surface. For example, due to diffusion the flow front in a tube is smeared out in the direction of its fore-axis, although to a minor extent compared to that caused by diffusion perpendicular to the axis. To understand the importance of the latter, one has to be familiar with the concept of the diffusion boundary layer.

As the net result of friction between the fluid and the wall of the tubing, the laminar flow rate profile is parabolic inside the tubing and the cuvette (cf. Eq. (2) and Fig. 1). In the vicinity of the walls convection is negligible relative to diffusion (i.e. the dimensionless Péclet number, defined as the ratio of advective transport rate to the diffusive transport rate [57], is \( Pe = 2 \pi R / D \ll 1 \)), but the former dominates further away from the surface (i.e. \( Pe \gg 1 \)). The zone in which diffusion is the dominant process is called the diffusion boundary layer; it has a thickness of [17,41,51]:

\[ \delta = \left( \frac{3}{2} \right)^{2/3} \left( \frac{D \cdot C}{v_0} \right)^{1/3}. \]  

(5)

where \( v_0 \) is the velocity of the leading point. \( \delta \) depends on the geometry of the fluidic system mainly through the constant \( C \). For example, in the case of a cylindrical cuvette with radius \( R \), \( C = 2R \), where \( z \) is the distance between the inlet of the cuvette and the point of sensing. It can be calculated from Eq. (5) that under typical experimental conditions \( \delta \) is 1–100 \( \mu m \).

2.3. Transport to, adsorption on, and desorption from the sensor surface

Surface-sensitive optical biosensors detect refractive index changes close to a solid–liquid interface (Section 3.1). Since the adsorption of the investigated objects (molecules, viruses, bacteria, vesicles, etc.) onto the sensor and their desorption from it can strongly affect the refractive index in the sensing zone, an adsorption model has to be considered for a complete description of detectable refractive index changes.

2.3.1. Adsorption model for single-component solutions

The simplest kinetic adsorption model of practical use distinguishes reversible (subscript \( r \)) and irreversible (subscript \( i \)) adsorption from a solution containing only a single type of analyte. To a first approximation the evolution of the surface excess of the reversibly and irreversibly adsorbing particles can be given by the following set of differential equations

\[ \frac{d \Gamma_r}{d \Gamma} = k_r c_v \phi(\Gamma) - k_d \Gamma_r \]  

and \[ \frac{d \Gamma_i}{d \Gamma} = k_i c_v \phi(\Gamma) \]

(6)

where \( \phi \) is the available area function characterizing the free surface area available to the adsorbing objects. Here, \( k_r \) and \( k_i \) are the rate coefficients for reversible and irreversible adsorption, while \( k_d \) is the first order rate coefficient for desorption; their actual values are characteristic of both the adsorbing object and the surface. The concentration of adsorbing particles at the bottom of the diffusion boundary layer is denoted as \( c_v \) (note that the layer probed by the sensor is generally much thinner than the diffusion boundary layer). The above model assumes that the reversibly and irreversibly adsorbing particles have the same size, shape, and surface affinity and that they randomly compete with each other. For example, a spherical, but chemically inhomogeneous protein could adsorb reversibly in certain orientations, otherwise irreversibly; the probabilities of the different orientations are subsumed in the rate coefficients. The total mass at the surface is the sum of reversibly and irreversibly adsorbed molecules, \( \Gamma = \Gamma_r + \Gamma_i \). Note that \( \phi(\Gamma) = 1 \) if \( \Gamma = 0 \), and \( \phi(\Gamma) = 0 \) if the whole surface is covered to an extent that there is not enough free space for further adsorbing objects. The available area function depends on the shape of the adsorbing object [58], and usually has a complicated polynomial dependence on the occupied surface area. Note that in the special case, when relatively large receptors are deposited on the sensor surface and used to capture a smaller analyte, the available area function has a simple linear form (Langmuir adsorption [59]).

A major aim of kinetic data analysis is the determination of the rate coefficients, which can be used to elucidate the molecular mechanism of an adsorption process. A key issue here is the appropriate characterization of the temporal evolution of \( c_v \), which is not always the same as that of the bulk concentration \( c_0 \).
The adsorption model described by Eq. (6) predicts that $d\Gamma/dt$ obtains its maximum value at the beginning of the adsorption process ($\Phi = 1$) and monotonically decreases as the desorption proceeds (Fig. 2 dashed line). In contrast, experimental data typically reveal a different behavior: at the initial stage of adsorption, a transient regime can be observed. The process is said to be transport-limited if the adsorption of the particle of interest is faster than the replenishment of material in the vicinity of the surface [60]. As adsorption proceeds, more of the surface is occupied by adsorbed molecules leaving less space available for newly arriving ones, resulting in a slower rate of surface mass increase. In this so-called adsorption-limited regime the rate of adsorption, $d\Gamma/dt$ decreases monotonically with time, reflecting the diminishing available area.

During transport-limited adsorption, $c_0$ is not equal to $c_0^*$; differences are the net result of diffusion from the bulk to the probe layer, and adsorption to and desorption from the surface. These can be taken into account by the following equation [40]:

$$V \frac{dc_v}{dt} = S \left( D \frac{c_0 - c_v}{\delta} -(k_i + \Gamma_i)c_v \Phi + k_d \Gamma_i \right),$$

where $V$ and $S$ are the unit volume and area, respectively. In a flow-through cuvette $c_0$ is kept constant above the diffusion boundary layer by supplying a continuous flow of analyte. If, however, flushing the volume above the diffusion boundary layer is not effectively instantaneous (compared to the sampling rate of the instrument and the duration of the whole adsorption process), a transient regime might appear in the kinetics of adsorption. As a result of this so-called flushing effect, $c_0$ and, therefore, $c_0^*$ are ill-defined in the early stages of solution exchange. The flushing effect can be aggravated by diffusion to and from the dead volumes — those volumes that are not part of the user-generated laminar flow and which, therefore, cannot be completely and effectively instantaneously flushed. Dead volumes act like reservoirs when a sample is exchanged in the cuvette and, therefore, initiate diffusion-driven changes in the local sample concentration; this essentially affects the OWLS signal, which will be considered later (Section 4.2.2).

### 2.3.2 Desorption from the sensor surface — readsorption and rebinding

If adsorption is irreversible, or to test whether it is, the adsorption stage with analyte in the fluidic system is followed by a washing/flushing stage when analyte-free medium (e.g., pure buffer) is introduced into the cuvette. During this stage $c_0$ (Eq. (7)) becomes zero and, typically desorption dominates (i.e., the term with $k_d$ becomes the most significant). It is important to note that $c_v$ is not zero except near the end of the stage.

Readsorption of freshly desorbed material occurs and is fully taken into account by using the same equations (6) and (7) for fitting the desorption part of the kinetics as for the adsorption part; the only difference is that $c_0$ is set to zero (exactly how this is done depends on consideration of the dead volumes, see Section 4.2.2.1).

It has to be noted that the above effect is also important in bioaffinity measurements, when receptors are immobilized on the sensor surface and ligands in solution are captured; if diffusion is slow compared to the on-rate (i.e., the dissociation phase is transport-limited), a considerably amount of ligand will rebind rather than diffusing into the bulk solution [60,61]. If this rebinding effect is not taken into account during data analysis, an off-rate that is much lower than the actual one will be obtained [59]. The transport-limited dissociation phase under flow conditions can be described, to a good approximation, with the special analytical formula developed by Schuck and Minton [62]. An alternative possibility is to prevent rebinding by adding competing ligand during the dissociation phase [59].

#### 2.3.3 Adsorption from complex, multicomponent analyte solutions

Importantly, the description of surface adsorption given in Sections 2.3.1–2.3.2 can be applied for single-component solutions only, and cannot be straightforwardly generalized for adsorption from a complex, multicomponent analyte solution. Adsorption from a complex sample takes place in a sequential and competitive manner, so the composition of the adsorbed layer is constantly evolving (Vroman effect) [63–66]. Those proteins which are bigger and have more affinity to the surface eventually displace the ones that occupy the surface more rapidly. As a result, for example the highest ratio of fibrogen to total protein at the occupied surface is generally obtained at intermediate blood serum concentrations [64].

### 2.4 Fluid handling in cell-based assays

Surface-sensitive biosensors enable various activities of adherent cells to be monitored with high sensitivity and high temporal resolution [14,30,67,68]. Cell-based studies using label-free biosensors can generally be classified as adhesion, signaling, or mechanotransduction assays, each of which requires different sample handling strategies. In general, sample handling in cell-based assays is always challenging. Additional difficulties with cell handling as compared to biochemical assays comes from that fact that living cells are highly dynamic entities very sensitively responding to changes in their surroundings, and are also able to actively modify their environment (e.g., through metabolism or secretion). Therefore, several criteria regarding sample handling have to be satisfied if one wishes to reliably monitor the behavior of living cells using OWLS or other surface-sensitive biosensors.

#### 2.4.1 Adhesion assays: general considerations, and cell deposition in an environment without continuous flow

In vivo, cell adhesion and spreading are induced by amino acid sequences presented by proteins of the extracellular matrix (ECM), which are specifically recognized by the adhesion receptors of the cell [69,70]. Once the connection between some adhesion receptors and ligands are established, the cell will begin to spread by actively reorganizing its cytoskeleton; this adhesion (attachment) is essential for most cells to survive and properly function. In most cases, therefore, the sensor should be precoated with either a synthetic (e.g., PLL–PEG–RGD [14,71]) or a purified (e.g., collagen, fibrinogen) polymer mimicking the ECM in order to specifically promote the spreading of the subsequently seeded cells.

In cell spreading assays, the molecular coating should ideally be able to specifically activate the adhesion receptors and repel soluble biomolecules at the same time; so any change in the signal will originate from the spreading of cells, and the adsorption of biomolecules secreted by the cells will not significantly contribute to it [14,71]. (Note that kinetic readout of multiple parameters (the position and width of the...
resonance peaks) in OWLS enables the refractive index variations caused by secreted molecules to be distinguished from refractive index changes provoked by cellular spreading [29,30,72]. However, this is a unique feature not common to all surface-sensitive biosensors.) The biosensor baseline is established with assay buffer (physiological buffer, in which the cells will be eventually bathed) above the precoated sensors. When seeding cells, a number of practical considerations should be taken into account. First, the introduction and deposition of the cells should ideally result in their uniform distribution on the sensor surface. Second, a sensing area coverage of somewhat more than 50% (typically meaning 2000–6000 deposited eukaryotic cells/mm²) is necessary to allow the relative adhesion strength to be determined; in OWLS-based cytometry, the refractive index change in the evanescent field at 50% coverage is a quantitative measure of cell-substratum adhesivity [46,67,73]. However, there is an upper limit to the desirable cell density; in order to allow uniformly distributed cells to have enough space for all of them to spread the formation of multilayers or that of cell aggregates should be avoided. Third, the duration of the deposition should be as short as possible; if there is significant cell-to-cell variation in the deposition time it is difficult to distinguish between signals corresponding to deposition and to the active process of spreading, since while some cells are still in suspension, others already on the surface will have begun to spread. One can try to separate the signals corresponding to the two processes via data processing — a number of numerical and theoretical models aim at describing the deposition kinetics of spherical particles sedimenting under the combined influence of Brownian collisions with the molecules of the bathing medium and the gravitational force [74,75]. However, it is more straightforward to achieve a quasi-instantaneous (compared to the time required for initiation of active spreading) deposition of cells. Cells in solution can be considered as spherical objects to an excellent approximation. Let us first assume them to have zero initial velocity. If they fall in a viscous fluid by their own weight, then terminal velocity is reached when the frictional force combined with the buoyant force exactly balances the gravitational force. The resulting terminal or settling velocity is given by [76,77]

\[ v_{\text{term}} = \frac{2(\rho_s - \rho_m)g}{\mu}r. \]  

(8)

The dynamic viscosity and the density of the aqueous medium can usually be assumed to be the same as that of the water, i.e., \( \mu = 6.92 \times 10^{-3} \text{ g/(cm} \cdot \text{s)} \) and \( \rho_m = 0.9933 \text{ g/cm}^3 \) at \( T = 37 \degree \text{C} \) respectively; the radius of a typical mammalian cell is \( r = 7 \mu \text{m} \) [78], its density is \( \rho_s = 1.1 \text{ g/cm}^3 \) [77]; and the gravitational constant is \( g = 981 \text{ cm/s}^2 \). This yields a terminal velocity of \( v_{\text{term}} = 16 \mu \text{m/s} \), which is so slow that concerns about the achievability of quasi-instantaneous deposition of all cells can be rightfully raised. However, cells can be made to move with velocity \( v_D \) during their introduction into the cuvette if the pipette with which they are introduced is oriented perpendicular to the sensor surface (forced deposition). Supposing that a volume of 0.1 ml suspension is run through the end of a cylindrical pipette tip having a diameter of 0.5 mm in a time interval of 5 s: then \( v_D = 10.2 \text{ cm/s} \), which is sufficiently high to result in the desired quasi-instantaneous deposition of all cells. This was confirmed experimentally (Section 4.1), when we used the forced deposition technique to seed cells on the sensor surface, and then successfully monitored their subsequent spreading. Forced deposition can be also achieved by centrifugation [79].

2.4.2. Mechatrontransduction assays: cells in a flow environment

Cells are responsive to mechanical stimuli; external forces influence the formation of adhesion sites, cell orientation, gene expression, and more [56,80]. To study the effects of external forces on cell behavior, cells are often investigated in a flow environment, an approach which may have more biological relevance as compared to experiments performed in the total absence of flow. The workflow of a standard mechatrontransduction assay starts with a surface functionalization step to obtain a cell-adhesive surface; then cells are seeded (preferably at high densities) and allowed to establish the first contacts with the surface for a given interval; and finally, a laminar flow with a well-defined rate is applied, which may be increased in a stepwise manner. However, increased scrutiny has to be given to the experimental design to avoid cell damage in such a study [81]. Cells in a flow environment may experience various forces; the most likely sources of cell injury are relatively large shear forces, interaction with small eddies occurring in turbulent flow, or interaction with bubbles during their breakup [81]; the last two can be easily avoided in a biosensor experiment, but the first one needs further consideration. In vivo exposure of various cell types (including red blood cells, immune cells, tumor cells, and endothelial cells) to flow predominantly occurs in the circulation system. The highest flow shear stresses are experienced in the arterial circulation, where time-averaged values are approximately in the range of \( 4 - 30 \text{ dyn cm}^{-2} \) [82]. In comparison, the wall shear stress (which is the maximum shear stress) of a Newtonian fluid during laminar flow in a straight cylindrical tube is [82]:

\[ \tau_w = \frac{4F}{\pi R^2}. \]  

(9)

In a typical biosensor experiment \( R = 0.0255 \text{ cm}, \) and \( F = 0.01 \text{ cm}^3/\text{s} \) (and \( F \) is the same as suggested for Eq. 8), which yields a wall shear stress of \( \tau_w = 5.3 \text{ dyn cm}^{-2} \); therefore no cell damage is expected for cells that tolerate the shear stresses in the circulatory system.

Molecular interactions can also be regulated by external forces. Flow-induced shear stress may provoke conformational changes in dissolved or adsorbed polymers [83], potentially leading to the exposure of otherwise buried molecular sequences (cryptic sites) [84], and/or to reduced or enhanced molecular interaction lifetimes (the latter interactions are called catch bonds) [85]). Consequently, a molecular surface coating that has been exposed to flow may promote cell spreading to a different extent than a coating that has not [84].

2.4.3. Signaling assays: the challenge of substance addition to pre-attached cells

Among all types of cell-based experiments carried out by means of label-free biosensors, probably signaling assays demand the most thought out sample handling strategy. One reason is that signaling assays inherently require high-throughput measurements and, therefore, a sample handling strategy that is compatible with high-throughput. Generally, the sensors are rendered cell-adhesive by creating a molecular coating on their surface via adsorption. Then cells are seeded at high density on the surface, and incubated for 1–2 days at 37 °C, 5% CO₂ to obtain a confluent cellular monolayer. These steps can be carried out manually with a channel pipettor. Subsequently, a baseline is established with assay buffer above the cells and some substance is added to them to activate a signal transduction pathway. Importantly, receptor activation and subsequent signaling are detected through both horizontal and vertical dynamic mass redistribution (DMR) in the bottom region of the cells [13]. Note that small changes, which cannot be revealed using conventional optical microscopy, are detectable in this way. The challenge lies in how the substance is introduced above the cells, especially because DMR signals are rather small and may change rapidly [79,86]. Ideally, introduction should be highly reproducible, and the flow generated during introduction should not perturb the cells (compare with Section 2.4.2). Obviously, these criteria are hardly met if substances are added manually with a pipettor; the introduction rate and the distance from cells are not well-defined and it cannot be done parallel in 96 or 384 channels. In closed cuvette systems (Section 4.1), therefore, primarily an integrated sample dispenser robot has been used for sample manipulation, which enables highly controlled and reproducible liquid exchange above cells (for a detailed protocol, see [79]). Cell signaling assays are considerably more
straightforward with flow-through fluidic systems [87]; here, cells should be plated on the precoated sensor surface, allowed to spread for a defined time interval, then a constant flow of buffer should be initialized and sustained throughout the assay. After baseline reading, any substance can be conveniently introduced into the flow. In addition, this system also enables the duration of the stimulation to be controlled. However, high-throughput biosensors with a flow-through system have not hitherto been commercialized.

3. Experimental: materials and methods

3.1. Optical waveguide lightmode spectroscopy

Surface-sensitive optical biosensors take advantage of surface-bound electromagnetic waves to detect refractive index changes (evoked by either bulk refractive index changes, molecular adsorption, cell spreading, or dynamic mass redistribution in spread cells) close to a solid–liquid interface. Exactly how this is done is greatly dependent on the technique itself and the actual instrumental configuration. OWLS detection is based on a nanograting planar optical waveguide (i.e., sensor chip, Fig. 3b), which is illuminated by a laser beam. The nanograting embedded into the waveguide structure enables light to be incoupled into the waveguide layer. The light then propagates several millimeters, permitting intensity measurements at the ends of the sensor chip. Such waveguiding, however, can be achieved only at discrete illuminating angles, which are dependent on the refractive index of the sample layer closest (up to 100–200 nm) to the surface of the sensor chip. The illuminating angle is varied by rotating the waveguide with a high-precision goniometer relative to the illuminating light beam (Fig. 1). Plotting the photointensity measured at the ends of the waveguide against the illuminating angle yield the OWLS spectrum; sharp resonant peaks with a typical width of 0.05–0.07° indicate at what angles waveguiding is achieved (resonant angles). Whenever the refractive index over the sensor surface is altered, the position of the peaks in the spectrum will be shifted.

Throughout this study, experimental data are presented as the alteration of the effective refractive index of the zeroth order transverse magnetic lightmode (simply denoted as ΔN). The effective refractive indices of the waveguide modes can be derived from the resonant angles [16,88].

3.2. Sensor chip preparation

OW2400 OWLS sensor chips (Microvacuum Ltd., Hungary) were used in all experiments presented in this study. Sensor chips were cleaned according to the following protocol. Cellular contamination was first removed by sonicating the chips in an aqueous medium. The waveguides were then soaked in chromic acid for 3 min, then rinsed with Milli-Q water (MQ), 0.5 M potassium hydroxide, and washed with copious amounts of MQ. The chips were then placed into MQ in a sonicator for 30 min and the water was changed every 3 min. Prior to experiments the waveguides were equilibrated in buffer overnight.

3.3. Experiments on flow-through systems

The prepared waveguides were mounted onto the measuring head of an OWLS instrument. Custom-made polyether ether ketone (PEEK) cuvettes were then sealed to the waveguide with a Kalrez O-ring [89]. Flow was guided by tubes made of either silicone (Ismatec, Tygon R3607) or polytetrafluoroethylene (PTFE), with inner diameters of 0.51 mm or 0.8 mm. The ends of the PTFE tubes were flattened with an

Fig. 3. Closed cuvettes in biosensorics. a) Image of a closed cuvette (shown both in assembled and disassembled form): i) cuvette and metal chip holder, ii) waveguide (highlighted in red), iii) side wall of the cuvette, iv) retaining screw. b) Closed cuvette used in OWLS experiments. i–iii) Schematic representation of the functional parts of an OWLS waveguiding sensor chip: supporting substrate, waveguiding film, grating incoupler, respectively. iv) Closed cuvette sealed by an O-ring to the surface of the waveguide forms the sample volume. v) Culture medium with which the OWLS baseline is established before monitoring of cell spreading. The cell suspension is introduced into the closed cuvette manually using a pipette. vi) OWLS signal obtained by monitoring cell spreading. The waveguide was coated with PLL (150 μl 0.1% solution incubated on the surface for 15 min at room temperature, then washed) and subsequently placed into DMEM buffered with 25 mM HEPES (pH 7.0). A suspension containing 20,000 3T3 cells was introduced into the closed cuvette with a pipette using forced deposition and subsequent spreading was monitored. The inset is a microscope image depicting the spread state characteristic of the cell line.
Omnifit kit (Biochem Fluidics) and connected via linear junctions. A porous hydrophobic membrane-based bubble trap (Omnifit) was integrated into the flow-through fluidic system.

When a large amount of sample was available (>2 ml), either a peristaltic pump (Reglo Digital, Ismatec) or a laboratory-built, computer-controlled syringe pump was used to generate continuous flow above the sensing area. In contrast, when the sample volume was limited, small amounts were injected into the fluidic setup either using an injection valve (i.e. SIS-06, see Section 4.3.1) or a septum injector (see Section 4.3.2).

For practical reasons, glycerol (Spektrum 3D) solutions were used as the sample in most OWLS experiments. Most importantly, the interaction of glycerol with the waveguide is completely and instantly reversible, i.e. the solution can be removed by flushing the fluidic system with MQ or PBS (phosphate-buffered saline, Sigma-Aldrich), resulting in restoration of the baseline. Hence, multiple experiments can be carried out consecutively with the same sensor chip. This is because glycerol only changes the bulk (cover) refractive index, i.e. it does not form an adlayer nor does it diffuse into the chip. For surface adsorption experiments the k-epsilon and k-omega formulations. The volumetric was applied to cope with the turbulence, which blends between the tum equations and the continuity equations were solved in a steady, near-wall boundary layer cells, which were hexahedral. The momen-

4.1. Closed cuvette without flow and manual fluid introduction using a pipette: monitoring cell adhesion and spreading

The simplest possible fluidic tool enabling the exposure of the sensing area to the solution of interest is the closed cuvette (Fig. 3). Samples have to be introduced manually using pipettes, and continuous flow cannot be generated in such an arrangement. However, all of the commercialized high-throughput optical biosensors employ open wells, and are not currently available with flow-through systems [14, 68].

Typically, closed cuvettes are used when some activity (adhesion and spreading, proliferation, response to effector molecules, etc.) of cells is monitored [14,25,26,28,30]. Depending on the aim of the investigation, cellular assays on a biosensor may take up to hours or days, and the fewer disturbances to the system during the measurement the better. Contamination can easily be avoided by covering the cuvette with a piece of Parafilm, but several undesirable phenomena can still potentially perturb the system. Diffusion of gases into the cell suspension can cause pH changes and solvent evaporation may cause the osmolality of the medium to increase. The biosensor might directly respond to such changes, which furthermore stress the cells, changing their normal behavior (i.e. that observable in an optimal, well defined and unchanging environment). Recently, an OWLS closed cuvette has been developed into a mini-inocubator that enables the temperature and pH of the cell suspension to be automatically controlled [30]. This mini-inocubator-equipped OWLS system has been used to monitor the spreading and adhesion of sensitive primary immune cells isolated from human blood [30].

The obvious caveat associated with the use of a closed cuvette system is the difficulty of performing manipulations on the sample. Nowadays one generally wishes to continue observation after the cells spread on the sensor surface and monitor either their proliferation, survival, or response to various effector molecules (drugs, ligands, toxins). It is known that not only the presence of the effector but also the duration of the stimulation is crucial in cell biology [90]. In contrast to desirable fast, yet gentle and controllable sample exchange, cumbersome pipetting from a closed cuvette implies the relatively uncontrolled removal of only a portion of the bathing medium. In addition, when the cuvette is mounted on the rotating goniometer of an integrated optical scanner (incoupling configuration [16]) the scanning has to be stopped to perform any manipulation in the sample volume of the closed cuvette and, therefore, typically for tens of seconds following sample addition the response cannot be monitored. In an OWLS device without moving parts, such as the outcoupling configuration [17] or one of the various kinds of interferometry [7,17], the measurement would not have to be stopped, although there might be some optical perturbation due to fluid movements. Furthermore, washing steps cannot be conveniently carried out in the closed cuvette. To overcome its drawbacks, flow-through systems for living cell applications have been specifically designed [77,87,90].

In summary, closed cuvettes are ideal for applications where sample manipulations during the measurement are rarely needed — these applications include cell spreading assays, or drug screening assays that aim at demonstrating a drug effect on a cell population where the duration of stimulation is less important (Table 1). A typical experimental arrangement with a closed cuvette and a spreading curve obtained with OWLS are shown in Fig. 3. Here, fibroblast cells were seeded onto the sensor surface precoated with PLL, and their spreading was monitored for 4 h.
Table 1
Summary of advantages and disadvantages of fluidic system components.

<table>
<thead>
<tr>
<th>Cuvette types</th>
<th>Flow-through cuvettes (Fig 1, Fig. 4/a,b,c,e,f)</th>
<th>Injector systems</th>
</tr>
</thead>
<tbody>
<tr>
<td>Closed cuvette (Fig.3)</td>
<td>Peristaltic pump</td>
<td>Syringe pump</td>
</tr>
<tr>
<td>Advantages</td>
<td>- Continuous washing</td>
<td>- Fully automated measurements</td>
</tr>
<tr>
<td></td>
<td>- Additional fluidic elements can be integrated</td>
<td>- Whole tubing can be made of PTFE</td>
</tr>
<tr>
<td></td>
<td>- May have more biological significance</td>
<td></td>
</tr>
<tr>
<td>Disadvantages</td>
<td>- Dead volumes act like reservoirs when samples are changed</td>
<td>- Cell suspension cannot be reproducibly introduced</td>
</tr>
<tr>
<td></td>
<td>- Complexity: all elements have to be tested carefully one by one</td>
<td>- Long tube lengths: diffusion has a significant effect when samples are changed</td>
</tr>
<tr>
<td></td>
<td>- Air bubbles can remain/appear between junctions in the system</td>
<td></td>
</tr>
<tr>
<td></td>
<td>- Temperature and pH have to be the same for all subsequently introduced samples</td>
<td></td>
</tr>
<tr>
<td>Typical applications</td>
<td>Monitoring cell adhesion and spreading</td>
<td>All kinds of biosensor experiments can be performed and monitored: protein adsorption, ligand-receptor binding, protein–lipid bilayer interactions, protein–DNA interactions, biocompatibility studies, cell response studies</td>
</tr>
</tbody>
</table>

4.2. Fluidic systems with flow-through cuvettes

Sophisticated fluidic setups can be built by connecting supplementary fluidic elements (bubble trap, junctions, pumps, etc.) to a flow-through cuvette (Fig. 1). Typically, the inlet tube connects the sample reservoir with the sample volume of the cuvette, and the outlet tube leads to a waste container (Figs. 1, 4). Peristaltic or syringe pumps are used to generate flow ensuring a constant supply of material. Exchanging the samples is very simple and the flow rate can be easily set to the desired value — altogether the flow-through technique is particularly advantageous because of the experimental controllability and simplicity it offers [17,41]. Moreover, since the flow is continuous, OWLS data is easily recorded during both the adsorption and desorption phases of a molecular/cellular process under well-controlled conditions. Therefore, the kinetic rate coefficients of the processes can be easily determined by fitting kinetic models of more or less sophistication to the data. The role of flow from a biological point of view was considered in Section 2.4.2; some investigations may have more biological significance if performed under flow, but this may substantially complicate data interpretation and/or experimental design.

Flow-through fluidics can, however, only be used when a sample amount sufficient for an entire experiment is available. With a typical flow rate in the range of microliters per second this usually means milliliters of solutions. Furthermore, given that OWLS (and DPI several orders of magnitude more so) is sensitive to changes in temperature and pH [47,89], it is critical to ensure that these parameters are the same for a subsequently introduced sample as those for the sample to be replaced. Moreover, some flow-through cuvettes might be inappropriate for working with mammalian cell suspensions due to geometrical issues; according to our experience, cells can adhere and aggregate in the immediate vicinity of the inlet aperture (before the sensing area), rather than being uniformly distributed on the entire bottom of the cuvette. Thus, careful scrutiny has to be given when designing a flow-through cuvette for cellular assays.

A flow-through system unavoidably hides risks in its relatively complicated arrangement and whole measurements can be endangered if the diverse constituent elements of the system are not carefully tested one by one and their incidental effects on the measurement revealed. Gas bubbles, for example, can grossly distort the data, but the integration of a bubble trap into the fluidic system offers an easy way to efficiently suppress this threat (Section 4.2.2.3). Also, the inner diameter of tubes has to be chosen carefully: it is advantageous to use larger diameters closer to the cuvette, followed by smaller diameters at the pumps to effectively dampen possibly abrupt pulsations/variations in flow.

The advantages, associated caveats and typical application areas of OWLS with flow-through fluidics are summarized in Table 1.

4.2.1. Basic building blocks of a flow-through setup

4.2.1.1. Cuvette and tubing materials. As previously found [89], and now confirmed, silicone cuvettes leave contamination on the surface of the
chip. We therefore used polyether ether ketone (PEEK) for the cuvettes and Kalrez (a perfluorinated elastomer) O-rings for sealing (Fig. 4), which are sufficiently inert to be recommended for all measurements.

Many materials, especially silicone, are unsuitable for the tubing. Those that are permeable may let gases diffuse into the sample, which may result in undesired bubble formation, especially when a peristaltic pump is used to generate flow (see Section 4.2.1.2). Tygon LF is soft enough to be suitable for peristalsis. PTFE (Teflon) is probably the best tubing material for connecting tubes, since it is resistant to most organic solvents and even to strongly acidic or basic solutions. However, PTFE is too rigid to be used with peristaltic pumping.

4.2.2. Exchange of samples using flow-through fluidics

Regardless of how carefully the samples are manipulated and exchanged, certain undesired effects cannot be eliminated and may significantly affect the recorded biosensor data, making their interpretation more difficult. Dead volumes are especially problematic (Section 2.2).

In contrast, a single peristaltic pump offers an easy method for generating quasi-uniform flow, but some commercially available peristaltic pumps (the ones with circular rather than elliptical roller races, or with fewer than 8–10 rollers) tend to introduce pulsations in the flow that can influence both the sensor itself[92] and the adsorption or other process under investigation.

### 4.2.2.1. Diffusion due to the dead volumes of the cuvette

Most flow-through cuvettes have unflushable volumes between the sealing O-ring and the inlet apertures. With properly planned experiments it can be shown that these “dead volumes” have an important effect on the actual biosensor measurements (see Fig. 5) – especially when small amounts of sample are used – and consequently these volumes should be minimized. As a result of diffusion to and from the dead volumes, data collection just after changing the flow from pure solvent to analyte solution may not represent the sample of interest.

Our test measurements were executed as follows. The baseline was established with either PBS or MQ, then pressure-driven flow of aqueous glycerol solution was initiated. The flow was suddenly stopped well before saturation of the OWLS signal, and a drastic decrease of the signal was observed (Fig. 5a). In the inverse experiment (Fig. 5b) the cuvette was initially fully filled with glycerol solution, which was then partially removed by pumping pure buffer for typically 0.5–1 min. In this case, stopping the buffer flow resulted in a rising biosensor
again and the same sequence was repeated.

The cuvette at the instant of stopping the flow is laminar and \( V_f \) is rather big (Fig. 6a). In contrast, if flow rates around 100 \( \mu l/s \) are used, the flow becomes turbulent and most of the dead volumes are successfully eliminated (Fig. 6b, and see Section 4.3.2).

4.2.2.2. Length of tubing. The Hagen–Poiseuille-equation (Eq. (2)) states that the flow resistance grows with tube elongation; one should, therefore, always check whether the original flow rate can be maintained if extra tubing is added. We used three tubes with lengths of 47, 147 and 447 cm and an inner diameter of 0.51 mm. The actual flow rate was determined from the amount of sample collected at the end of the tubing and the collection time. Our pump was robust enough for flow rate not to diminish with tube elongation.

An earlier investigation found no effect of tube elongation on the sensor signal saturation time [93]; however, tubes with only slightly different lengths were used in that study (17.3, 22.3 and 25.3 cm). In contrast, we found that significantly more sample is necessary for reaching saturation of the OWLS signal (i.e., to completely fill the cuvette with the sample) when the inlet tube is longer, and suspected that this was an effect of diffusion. It is clear that the diffusion in the direction perpendicular to the axis of the tube is much more significant than diffusion parallel to it [94]: sample in the boundary layer along the walls of the tube is exchanged by diffusion instead of convection.

The time the fluid spends in the tubing (i.e., the average time available for diffusion) is proportional to the tubing length \( L (t \propto L) \). The distance the sample diffuses perpendicular to the diffusion boundary layer is proportional to the square root of the time spent in the tubing (Eq. (3)), hence, the saturation time of the signal \( t_s \) is expected to increase proportionally to the diffusion time \( t \). Saturation times measured at different tubing lengths and plotted against the square root of tubing lengths can be nicely fitted with a straight line \( t_s \propto L^{1/2} \); thus the experimental data well supports the above prediction (Fig. 7).

4.2.2.3. Eliminating bubbles — effect of a bubble trap. A bubble forming inside the fluidic channel has a grossly different refractive index compared to the liquid medium or analyte and its presence will, therefore, severely distort the biosensor experiment — therefore great care has to be taken to avoid bubbles. Sonication, filtration and vacuum treatment of the solutions degases them and, hence, reduce the probability of bubble formation. Wider tubing followed by a narrower one helps to prevent bubbles forming at the junctions between tubes.

Another possibility is to incorporate a bubble trap into the fluidic setup [47]. Although the inner part of the bubble trap contains multiple arcs in which the sample is guided, we found that it has no undesirable mixing properties. Only a slight increase in \( t_s \) was observed when the bubble trap was integrated into each of the three tubes having different lengths (47, 147 and 447 cm) as compared to the cases when we did not use a bubble trap. The increase corresponded to an increased tubing length: a 15 cm long extra section of tubing was inserted to integrate the bubble trap, which itself contributes the equivalent of an additional 15 cm (the approximate length of its arcs). This is evidenced in Fig. 7, as these additional data points, marked with "bt", are well fitted by the model described in Section 4.2.2.2.

4.3. Injection systems for the introduction of limited amount of sample

Some samples are scarce or highly expensive and, therefore, only very limited amounts may be available. The minimum amount of sample necessary for an experiment (which normally means enough to obtain the kinetics up to steady state) can be effectively decreased if the sample is not pumped through the whole fluidic setup but injected closer to the sensor. It should be stressed, however, that small sample amounts are more prone to attenuation caused by diffusion (Sections 4.2.2.2, 4.3.1).

Our experimental findings were qualitatively confirmed by computational simulations of the flow in one of our flow-through cuvettes (Fig. 4a). When the flow rate is low (1 \( \mu l/s \)), the flow is laminar and \( V_f \) is rather big (Fig. 6a). In contrast, if flow rates around 100 \( \mu l/s \) are used, the flow becomes turbulent and most of the dead volumes are successfully eliminated (Fig. 6b, and see Section 4.3.2).

Signal, clearly indicating that glycerol was diffusing into the measuring zone from the unflushed volumes.

Here, we introduce a method to estimate the size of the dead volumes relative to the total cuvette volume. The effective refractive index \( N \) is approximately linearly proportional to the change in refractive index of the cover layer \( (n_c) \) [88], i.e.

\[
\Delta N = \left( \frac{\partial N}{\partial n_c} \right) \Delta n_c.
\]

Denoting the volume that is flushed with laminar flow (convective zone) at time \( t \) by \( V_C(t) \), and the corresponding unflushed dead volume (diffusive zone) by \( V_D(t) \), the total volume of the cuvette is

\[
V_T = V_C(t) + V_D(t).
\]

Let \( \Delta N_1 \) and \( \Delta N_2 \) be defined as in Fig. 5a, and \( \alpha \) be the constant of proportionality between the effective refractive index and the concentration of the sample. Using this notation, the amount of glycerol in the cuvette at the instant of stopping the flow is \( V_C(t) \alpha \Delta N_2 \) and the total amount of glycerol following equilibration of local concentration differences is \( V_f(t) \alpha \Delta N_1 \). Building on the fact that the amount of glycerol present in \( V_T \) does not change after the flow is stopped, the two quantities can be equated, yielding

\[
V_D = V_f \left( \frac{\Delta N_1 - \Delta N_2}{\Delta N_2} \right).
\]

Fig. 5. Estimating the unflushed (dead) volume of a flow-through cuvette. a) Before \( t = 0 \) PBS was pumped at 1.4 \( \mu l/s \). At \( t = 0 \) (marked with an upward pointing arrowhead), the flow was changed to a 6% solution of glycerol in PBS. At \( t = 1 \) min (marked with an arrowhead) the flow was stopped. At \( t = 3.5 \) min (marked with an arrowhead) flow of glycerol was resumed until \( t = 9 \) min, then changed back to PBS. For further explanation see the text. b) The inverse experiment. Before \( t = 0 \) the cuvette was completely filled with glycerol. At \( t = 0 \) PBS was pumped until the point marked with \( \perp \), PBS flow was resumed at the next arrowhead. At the end of the cycle the cuvette was completely refilled with glycerol again and the same sequence was repeated.
A combination of the reproducible and precise fluid handling characteristic of flow injection analysis (FIA) with the sensitive optical detection offered by OWLS resulted in the development of FIA-OWLS immunoassays. An injection valve (Section 4.3.2) and manual injections have been used to introduce samples into the cuvette [95]. For the development of OWLS immunosensors an FIA system with a peristaltic pump, an injector valve and an injector loop have been employed [96, 97]. Various lab-built flow-through cuvettes and injection ports have been tested to reproducibly introduce small amounts of samples. In polyelectrolyte studies normally 100 μl sample solution was manually introduced to flush a 37 μl cuvette [98–100]. Elsewhere, a sequence of adsorption steps has been used to maximize adsorption from a given sample quantity [101–103]. Very recently, the sophisticated FastStep™ [104] and OneStep™ [105] injection systems have been developed. In the former configuration the sample and buffer streams merge right before the flow cell, and a stepped analyte concentration profile is produced by varying the flow rate ratio of the two branches. In the latter configuration, the undiluted analyte disperses in a capillary tube before entering the cuvette, thereby producing a concentration gradient. In contrast to standard fixed concentration injections, where a set of dilutions is required to complete a dose-response range, both of these systems generate a full dose response from a single analyte injection, thereby reducing the variability, hence systematic error, among experiments. Therefore, these systems enable a gradient in pH, salt, or co-factors for rapid optimization of buffer conditions to be conveniently titrated.

In this paper the SIS-06 injection valve supplied to a BIO-210 OWLS instrument and a septum injector are tested and discussed in more detail.

4.3.1. Injection valves

The SIS-06 injection valve (Fig. 8) from MicroVacuum Ltd. can optionally be integrated between a pump and a regular flow-through cuvette to reduce the amount of sample necessary for a measurement. Both the buffer and the limited amount of sample are transferred into the cuvette by continuous pump-driven flow. The injector has 6 channels, and the exact route of flow depends on the operation mode set. The injection valve operates in two modes. In “Load” the sample

---

**Fig. 6.** Results of computational fluid dynamic simulation modeling flow in our cuvette (Fig. 4a) at a) low (1 μl/s) and b) higher (100 μl/s) flow rates. Explanation of colors: the red volume is moving with at least 1 mm/s, while the blue volume is considered as stationary. The color bar represents the lifetime of streamlines (represented as individual thin lines). Streamlines in the figures suggest that flow in the cuvette is laminar (there are no currents perpendicular to the direction of flow, nor eddies or swirls of fluid) at a flow rate of 1 μl/s, and turbulent at 100 μl/s. In the former case, huge volumes remain unflushed in the cuvette (cuvette volumes in blue color, panel a); acting as dead volumes during sample exchange. Turbulent flow, on the other hand, enables effective sample exchange in the whole cuvette (nearly all blue volumes are eliminated, panel b).
loop can be filled with the analyte, while the buffer is conveniently moved towards the cuvette through two channels that bypass the sample loop. In “Inject” mode the sample loop is connected to the pressure-driven buffer flow and its content is transferred to the cuvette. Calibrated sample loops are commercially available with different volumes. Throughout this study we used one with a volume of 50 μl.

First we injected 5.8% glycerol into the fluidic setup via the SIS-06 injector valve (Fig. 9a). The glycerol sample (n = 1.33518) and MQ (n = 1.33085) was pumped over the sensor with a programmable syringe pump at different flow rates (1.44, 0.7 and 0.14 μl/s). Despite the fact that the volume of the cuvette was only about 20 μl, the injected 50 μl of sample seemed to be insufficient to flush the cuvette through, we could not saturate the signal using any of these flow rates (Fig. 9a). We suspected that this was a consequence of the presence of unflushable volumes in the tubes and the cuvette (Section 4.2.2.1).

The minimal sample amounts necessary to saturate the signal at each flow rate were determined in separate experiments, in which an unlimited amount of sample was pumped until a signal plateau was obtained (Fig. 9b). Surprisingly, more than 200 μl of sample was required when a flow rate of 1.4 μl/s was used, and only slightly less was needed if the flow rate was 0.14 μl/s. This underlines the importance of optimizing the fluidic arrangements before injecting small amount of samples, in order to obtain relevant results without wasting material.

In order to minimize diffusional effects and waste, the fluidic setup has been rearranged according to the conditions discussed in Section 4: we have i) changed tubes to ones having smaller inner diameters (from 0.8 mm to 0.51 mm); ii) designed a cuvette having a smaller volume (approximately 15 μl); iii) reduced the tubing length between the valve and cuvette to 4.5 cm; iv) chosen low flow rates; and v) integrated a bubble trap in the system. We then performed a new experiment with the biosensor and used the injection valve for sample introduction. Using this optimized fluidic system we were finally able to successfully measure the presence of 50 μl of 6% glycerol (Fig. 10), but only when the flow rate was at most 0.14 μl/s (as discussed earlier, a smaller amount of sample is enough to saturate the signal when a lower flow rate is used).

It should be realized, however, that the SIS-06 injector used with continuous flow is less suitable for adsorption or affinity-binding studies, because these processes generally take more time than for which the sample could possibly be present in the cuvette. Keeping the sample longer in the cuvette by lowering flow rate is strongly discouraged, because the binding process would become transport-limited. Stopping the flow when the cuvette is completely filled with the analyte solution may be a better option, but careful calibration is needed to determine...
molecular adsorption, receptor underlying the investigated surface process (whether cell spreading, turn, can contribute to a better understanding of the mechanisms

5. Conclusions

eliminates transport-limited retardation of the initial adsorption rate.

the exact time the flow has to be stopped (because protein solutions often do not alter the signal immediately).

4.3.2. Septum injector

A special cuvette equipped with a septum injector port (Fig. 4/d,e) enables limited amounts of sample to be directly injected over the sensor. To introduce the sample, a membrane on one of the “stalks” of the cuvette has to be pierced with a special septum needle; the withdrawal of the needle allows the membrane to self-seal.

With measurements using 50 μl of 6% glycerol (Fig. 11a) we demonstrated how the problem of dead volumes could be eliminated by a fast injection rate, which was sufficient to effectively flush the dead volumes (cf. Fig. 6). The adsorption of 50 μl of 50 μg/ml avidin rapidly introduced via the septum injector also gave satisfactory results (Fig. 11b). Desorption required multiple injections of pure buffer, which, however, generates complicated kinetics. Hence, it is desirable to flush buffer continuously (which is usually available in unlimited quantity) to obtain a monotonic desorption signal, which then can be analyzed to determine the kinetic parameters [62]. To achieve this, we removed the needle of a septum syringe and introduced it into a piece of tubing (Fig. 4/e), which could be readily used to connect the pump and the septum cuvette, thus enabling continuous washing (Fig. 12). This modification eliminates transport-limited retardation of the initial adsorption rate.

5. Conclusions

One of the most important advantages of label-free biosensors is that they generate kinetic data, thus allowing kinetic data analysis. This, in turn, can contribute to a better understanding of the mechanisms underlying the investigated surface process (whether cell spreading, molecular adsorption, receptor–ligand interactions, etc.) — which is ideally the final aim of every study in the field. It should be realized, however, that the way the liquid samples are handled (introduced, exchanged) may deeply influence both the kinetics of the surface process and the detection limit of the biosensor. The fluidic system is the most critical part of any biosensor setup, since the majority of experimental artifacts and misinterpretations of the data generally originate in an inappropriate fluid handling strategy or in the neglect of fluidic effects that severely interfere with the kinetics of the true surface process.

In an adsorption or receptor–ligand binding assay, for instance, the obtained biosensor signal will not be dominated by the kinetics of the true surface process, unless transport limitation of the analyte (Sections 2.1, 2.3.1) and the flushing effect (Section 2.3.1) are both successfully eliminated. Diffusion to the diffusion boundary layers (Section 2.2), or to the dead volumes (Sections 4.2.2.1, 4.3.1) of the cuvette may cause the analyte solution to be significantly attenuated, therefore the sample concentration in the sensing zone will be ill-defined, precluding the correct interpretation of the obtained kinetic data. (When adsorbing macromolecule solutions are measured, the effect of attenuation may be masked by adsorption; data analysis ignoring this distortion is likely to lead to serious error.) These phenomena, illustrated with experimental data and discussed in Section 4.2, have an increasingly decisive role when working with strongly limited amounts of samples (Section 4.3.1). These effects can be most successfully reduced by generating turbulent flow in a septum injector-based fluidic system (Section 4.3.2, Fig. 6); this efficiently eliminates those volumes that cannot be flushed with laminar flow. If necessary, the dead volumes can be further decreased if a bigger amount of solution is injected into the cuvette. However, efforts to minimize the necessary amount of sample for the actual measurement and to flush the cuvette perfectly are, to some degree, opposed to each other; consequently, one always has to make a compromise. If the sample of interest is very expensive and/or scarce, calibration with a model solution (e.g. glycerol), and subsequent correction of the signal of the actual sample can be the key for more accurate measurements. In the future, further miniaturization of the instrumentation may take place and, when molecular samples are being investigated, the tube dimensions may be further...
reduced down to the nanoscale. This introduces certain peculiarities that should then be explicitly considered[106,107].

Sample handling of cell suspensions may be even more challenging than that of analyte solutions (Section 2.4), especially because cells are capable of sensitively responding to changes in their surroundings and actively modifying their environment. Ideally, both the temperature and pH of the bathing medium should be kept at constant levels, preferably close to that experienced by cells in vivo; this requires further developments on a basic fluidic system. Interpretation of kinetic cellular data may be complicated because optical biosensors detect refractive index changes in a nonspecific manner (Section 3.1). Considering a cell spreading assay, the biosensor signal probed by cell spreading may be superseded by an adsorption signal if cellular secretion adsorbs or binds to the sensor surface or the molecular coating (Section 2.4.1). Monitoring other cellular activities may demand further considerations (Sections 2.4.2, 2.4.3). Appropriate flow-through fluidic systems enable cellular mechanotransduction to be monitored with a biosensor (Fig. 1), but this needs careful scrutiny in the case of other cell types (Section 2.4.2).

In summary, when measurements are carried out by means of biosensors, it is critical to establish a reliable strategy and a well-tested fluidic system (Section 4) for controllable and reproducible fluid handling. There seems to be no ideal fluidic design that is optimal for every application; each should be matched appropriately. In Section 4 we reviewed the most common fluidic systems and components used in biochemical surface sensing, illustrated their performance with experimental data, and discussed their advantages and disadvantages; our final conclusions are summarized in Table 1.

Acknowledgments

The support of the Hungarian Scientific Research Fund (OTKA-PD 73084) and the European Commission (ERG OPTIBIO) is gratefully acknowledged. Part of this work was carried out in the framework of a European Commission-supported project, aiming at blood analysis using optical biosensors (PSENS). This work was supported by the Lendület program of the Hungarian Academy of Sciences, and the Bolyai Scholarship to B. S. from the Hungarian Academy of Sciences.

This research was realized in the framework of TÁMOP 4.2.4. A-2/11-1-2012-0001 “National Excellence Program – Elaborating and Operating an Inland Student and Researcher Personal Support System Convergence Program”. The project was subsidized by the European Union and co-financed by the European Social Fund.

References

[10] Patko D, Cottier K, Hamori A, Horvath R. Opt Express 2012:20:23162–73. Finally, the cuvette was flushed with a continuous flow of PBS, PBS containing 0.05%Tween 20, and pure PBS again. For further experimental details, see[39]. Figure is adapted from Ref. [39].

Fig. 12. Experimental usage of the modified septum cuvette-based fluidic system. A septum needle was introduced into the end of a tube; this tube then could be readily introduced into the septum to connect the cuvette with a flow reservoir. This system enables a flexible exchange between two sample introduction strategies: continuous flow and injection of highly restricted sample amounts (Fig. 4e). Preceding the online experiment, the OWLS chip underwent an ex situ preparation procedure: its surface was functionalized with polyethylene imine (PEI) and biotinylated with NHS-biotin. The OWLS experiment was initialized by establishing a baseline, and then the prepared sensor surface was exposed to avidin by introducing a continuous flow of 50 μg/ml solution through the modified septum cuvette. After a washing step with PBS, the pump generating the analyte flow was stopped and 50 μl anti-CRP (C-reactive protein) antibody was introduced into the cuvette by manual injection. A ~40 min incubation period was terminated by flushing the cuvette with a continuous flow of PBS. Next, 50 μl of 10 μg/ml CRP solution was injected into the cuvette and allowed to bind to the surface for ~20 min. Finally, the cuvette was flushed with a continuous flow of PBS, PBS containing 0.05% Tween 20, and pure PBS again. For further experimental details, see[39]. Figure is adapted from Ref. [39].