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(71) Applicant (for all designated States except US): **NANO S BIOTECHNOLOGIE GMBH** [AT/AT]; Gregor Mendel-Strasse 33, Uni f. Bodenkultur/ZUF, A-1180 Vienna (AT).

(72) Inventors; and

(75) Inventors/Applicants (for US only): **SLEYTR, Uwe** [AT/AT]; Industriestrasse 78/26, A-1220 Vienna (AT). **PUM, Dietmar** [AT/AT]; Mittelgasse 19/14, A-1060 Vienna (AT). **GUFLER, Petra, Christine** [AT/AT]; Hasenauerstrasse 18/1/4, A-1190 Vienna (AT).

(74) Agent: **SONN & PARTNER PATENTANWÄLTE**; Riemergasse 14, A-1010 Vienna (AT).

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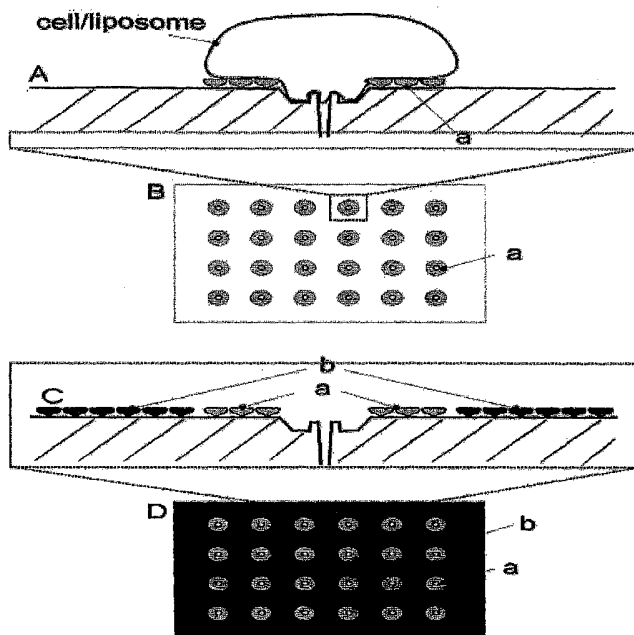
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(54) Title: SUBSTRATE FOR A DEVICE FOR ELECTROPHYSIOLOGICAL MEASUREMENTS



(57) Abstract: A substrate for a device for electrophysiological measurements comprising at least one perforation, wherein the substrate is covered at least in part with at least one protein and/or glycoprotein layer.

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Substrate for a device for electrophysiological measurements

This invention relates to a substrate for a device for electrophysiological measurements.

In recent years the patch-clamp technique became state of the art for electrophysiological measurements of ion channels and transporters located in cells, cell fragments or artificial lipid membranes. The classic patch-clamp technique uses a glass pipette, which is gently pressed against the lipid membrane. A very small area is isolated, which is investigated (Sakmann & Neher, 1995). The electric currents through reconstituted ion channels and pores are recorded in a highly specific and sensitive way. However, classic patch-clamping is time-consuming and not convenient for practical devices and automation. Recently, automated patch-clamp devices ("patch clamp on a chip" (Sigworth & Klemic, 2002)) were introduced, using planar microstructured chips made of PDMS (Klemic et al., 2002), silicon (Pantoja et al., 2001; Schmidt et al., 2000) or glass (Fertig et al., 2002; Brüggemann et al., 2003) for recording from whole cells (Fertig et al., 2002; Klemic et al., 2002), liposomes (Schmidt et al., 2000) and painted artificial membranes (Pantoja et al., 2001).

The planar chip replaces the pipette. Patch-clamp-on-a-chip enables simultaneous patch-clamp recordings from many liposomes or cells. The decisive factor in this system is the formation of a high-resistance gigaohmic seal between the chip and the lipid membrane. Lower resistances would lead to a low signal to noise ratio. In all systems difficulties with seal formation were observed, presumably because of non-sufficient contact area (narrow sidewalls, sharp corners of the etched aperture).

It is an object of the present invention to provide a substrate for a device for electrophysiological measurements which allows an accurate positioning of the membranous sample on the substrate and supports the formation of a high-resistance gigaohmic seal between a substrate and a membranous sample.

Therefore the present invention provides a substrate for a device for electrophysiological measurements comprising at least one perforation, wherein the substrate is covered at least in part with at least one protein and/or glycoprotein layer.

According to the present invention "the at least one perforation" can exhibit different three dimensional shapes. For in-

stance the perforation has the form of a frustum of a cone or a pyramid, of a cylinder, of a wedge, of a prism or of a cube.

According to the present invention "substrate" is defined as a structural component of a device for electrophysiological measurements whereon a membranous sample can adhere. Furthermore the substrate may function additionally as a barrier between two electrode compartments of a patch clamp device.

The "protein layer" according to this invention refers to all kind of proteins capable to be immobilised on the surface of a substrate to be used in apparatus for electrophysiological measurements. Examples for suitable proteins are S-layer proteins, antibodies, other proteins and glycoproteins and fragments thereof. Of course also polypeptides and peptides may be immobilised on the surface of a substrate.

The substrate according to the present invention may be used in apparatus for electrophysiological measurements employing the patch clamp technique. Such apparatus are disclosed for example in the WO 96/13721 A1, WO 99/66329 A1 and US 2002/0108869 A1. The presence of more than one perforation enables the substrate to be used also in high throughput screenings. Furthermore the invention enables standardized and reproducible spatial binding properties of membranous samples for electrophysiological measurements.

The protein layer (glycosylated or non-glycosylated) can be bound covalently or non-covalently, e.g. by hydrophobic or hydrophilic interactions, to the surface of the substrate. Furthermore the substrate can be covered entirely or partially with a protein layer, whereby the protein layer may comprise more than one protein species.

The at least one protein layer provides anchoring/adhesion sites for membranous samples, e.g. cells, cell fragments, liposomes and other lipid membrane aggregates, on a substrate which can be used in patch-clamp devices known in the state of the art. Furthermore the protein layer plays an important role in the high-resistance gigaohmic seal formation between the lipid membrane and the chip surface. Moreover the proteins on the surface of the substrate can be used as a pre-blocking inert coating, in order to prevent interfering effects and unspecific adsorption of cells, liposomes and other lipid aggregates. The invention enables a sensitive electrical acquisition of transmem-

brane proteins and is a further step toward high-throughput screening for drug discovery efforts.

According to a preferred embodiment the at least one protein layer is modified with at least one functionality. A modification of the protein layer with at least one functionality enables the substrate to interact with e.g. the membranous sample. The interaction of the at least one protein layer with the membranous sample is helpful in stabilizing the membrane on the substrate and furthermore to create a gigaohmic seal. The modification of the protein layer can occur chemically (by covalent or by non-covalent binding) or by genetic engineering (by creating e.g. fusion proteins).

The possibility to create a broad spectrum of recombinant proteins which can be employed in the formation of a protein layer (e.g. S-layer fusion proteins) enables specific interactions between the protein and specific groups (ligands, receptors) on the liposome, lipid bilayer or cell surface. Thus, a protein layer, in particular an S-layer, allows a tight and specific binding of unlimited types of liposomes, lipid bilayer aggregates, whole cells or cell fragments in an extremely gentle way. Thereby gigaohmic seal formation (with or without a gentle suction) is greatly facilitated.

The binding mechanisms and functionalities exhibited by the protein layer are the following (e.g. S-layer protein(s)):

- *) Lattice of native or recombinant S-layer proteins, or a mixture of native and recombinant S-layer proteins (co-recrystallization):
 - 1) unmodified S-layer lattice; utilization of the intrinsic binding affinities (non-covalent interactions).
 - 2) chemically modified S-layer lattice providing specific physico-chemical properties and binding affinities.
 - 3) Attachment of cells/liposomes/lipid aggregates due to topographical features of the S-layer lattice (edge sharpness of zone/chip boundaries and geometry).

- *) Reactive groups/receptors on the S-layer recognize ligands in the lipid membrane; reactive groups/receptors in the lipid membrane recognize an S-layer-bound ligand; e.g. antigen-antibody, ligand-receptor, lectine-carbo-

hydrate-interactions, metal-affinity-tags (e.g. histidine); S-layer/streptavidin fusion protein lattice - specific binding via biotinylated molecules in cells/liposomes/lipid aggregates.

According to another preferred embodiment the at least one functionality is selected from the group consisting of proteins, polypeptides, peptides, ligands, chemical groups and mixtures thereof. The functional groups of the at least one functionality may comprise all functional groups known in the state of the art, which can be used for protein modifications. For instance these groups comprise proteins (e.g. avidin, streptavidin, lectine, antibodies and fragments thereof), ligands (e.g. carbohydrates, biotin) and chemical groups (e.g. -OH, -COOH, -NH₂). For further details see Wong (1991) and Schalkhammer (2002).

Preferably the at least one protein layer is composed of at least one S-layer. The chemical and structural features of S-layer proteins turned out to be advantageous for a substrate to be used in patch clamp applications.

S-layer proteins are routinely utilised in several technical applications. For instance the EP 0 306 473 B1 discloses the use of S-layer proteins as a carrier for haptens, immunogenic or immunostimulant substances. The immobilisation of molecules, particularly of proteins, on S-layer proteins is described in the EP 0 362 339 B1. The WO 01/81425 A1 discloses the use of a structure comprising carbohydrates (secondary cell wall polymer) as an anchoring means for S-layer proteins in order to bind them to a solid support. In the WO 02/097118 A1 the production of a layer of functional molecules on a substrate using S-layer proteins is disclosed. Therein the protein layer on the surface of a carrier is formed by creating an electrochemical potential difference between the solution and the surface. The EP 0 189019 B1 discloses the making and the use of an ultra-filtration membrane employing S-layer proteins.

S-layer protein species from different organisms can be used. A protein species is used as a native, a recombinant, or a recombinant protein optionally with distinct functional domain(s), e.g. streptavidin, ligand, affinity tag. The S-layer is composed of identical S-layer proteins, a mixture of at least two species, or different protein species with each comprising

one or more specific functionalities. The ratio of the distinct proteins can be varied. The S-layer can be native or chemically modified. Furthermore S-layers can be recrystallized as mono-, bi- or multilayers.

Dependent on the S-layer protein species employed, distinct surface properties are obtained after recrystallization:

- *) intrinsic functionalities of native isolated S-layer proteins and recombinant S-layer proteins with no fused functional domain: e.g. charge density and distribution. In many cases the recombinant S-layer proteins that are predominantly expressed in *E. coli* show a different charge distribution than the native protein isolated from the bacterial cell wall.
With regard to their orientation in vivo, a common feature of S-layers is their smoother charge-neutral outer surface and a more corrugated, net-negatively charged inner surface (Pum et al., 2000). Hence, the orientation of the S-layer proteins on the substrate defines the physico-chemical properties.
- *) functionalities after chemical surface modification of native S-layers; e.g. change in charge and charge density, introduction of chemically reactive groups.
- *) artificially inserted functional domains : genetically engineered S-layer fusion proteins, chemically fused functional domains.
- *) inert surface character: neutrally charged surfaces. These types of S-layers show no binding affinity or unspecific binding.

Two-dimensional crystalline bacterial surface layers (S-layers) can be recrystallized in vitro into coherent lattices on a great variety of surfaces and interfaces, including silicon or glass (Sleytr et al., 2002, 2003; Pum et al., 2000). Lab-on-a-chip-technologies were facilitated with the patterning of S-layers on solid supports by microlithography (Neubauer et al., 1997, 1998; US 6296700 B1) or micromoulding in capillaries (Györvary et al., 2003a). Moreover, functional groups of the at least one functionality on S-layers interact with the lipid head group region of phospholipids, leading to a stabilization of

lipid membranes. Stabilization was demonstrated on lipid membranes (Schuster et al., 1999) (Wetzer et al., 1997, 1998; Gufler et al., 2004) and on liposomes (Küpcü et al., 1995; Mader et al., 1999). S-layers do not penetrate the hydrophobic region of mono- (e.g. tetraetherlipid) or bilayer membranes; therefore they have no impact on the integrity of the lipid aggregates, cells or cell membranes. S-layers provide a natural environment for lipid membranes and biological materials: e.g. in many archaeal organisms the S-layer is the only component external to the cytoplasmic membrane (Baumeister & Lembcke, 1992). A broad range of S-layer proteins with different physico-chemical surface properties and functionalities are available. The implementation of recombinant S-layer- (US 2002/0168728 A1) and S-layer-fusion proteins exhibiting functional domains, enables the application of specific binding mechanisms (Moll et al., 2002). Due to the high regularity of the crystal lattice, functionalities repetitively arranged in well defined positions and orientations can be obtained. Furthermore a significant reduction in surface roughness of the substrate is observable after recrystallization (Gufler et al., 2004).

According to a preferred embodiment of the present invention the substrate comprises an electrically insulating material. It is a basic requirement of devices for electrophysiological measurements, e.g. patch clamp devices, that the current can only pass the substrate through the pores, transporters and channels part of a membranous structure present in the vicinity of a perforation of the substrate. Therefore substrates used for patch clamp experiments may comprise electrically insulating material. The surface of the substrate or the entire substrate may comprise an electrically insulating material.

Preferably the electrically insulating material is selected from the group consisting of silicon, glass, solid state polymers and mixtures thereof. Microfabricated chips made of silicon, glass, solid state polymers or other materials are preferably used. Transparent chips, made e.g. of glass or synthetic polymers, can be used for simultaneous electrophysiological and optical or spectroscopical measurements (e.g. fluorescence measurements, microscopical methods). According to another embodiment part of the substrate may also comprise an electrode. In this case part of the substrate is electroconductive.

The perforation is preferably 100 nm to 10 μm , preferably 500 nm to 9 μm , more preferably 1 μm to 8 μm , particularly 2 μm to 7 μm wide. The perforations of the chips are fabricated e.g. by nano- and micro-lithographic methods, laser or micromoulding techniques. The dimension of the perforation is adjusted according to the size of the membranous sample, whereby different perforations of the substrate can possess differing dimensions. This demonstrates once more the flexible use of the substrate according to this invention.

"Wide" according to this invention refers to the physical dimensions of the perforations. For instance the term "wide" for a circular perforation refers to its diameter and "wide" for a quadratic perforation is referred to the distance between the nooks of such a perforation.

According to a preferred embodiment the substrate is substantially planar. The geometry of a substrate can be manifold. For instance the substrate can be planar like chips or exhibit a spherical form.

According to another preferred embodiment the substrate is chemically or physically modified. Not only the protein layer on the surface of the substrate can be modified, but also the surface of the substrate itself. Such modifications allow creating zones in which e.g. no protein layer can adhere and other zones where an interaction between surface of the substrate and the membranous sample is prevented. Other modifications may prevent an unspecific binding of the membranous sample on the surface of the substrate, where no protein layer is present.

Such a modification is preferably selected from the group consisting of ionizing radiation, atomic radicals, corona treatment, silane groups, chemical groups and mixtures thereof. Chemical groups may comprise carbohydrates (e.g. secondary cell wall polymer), biotin and reactive groups. Other chemical groups and modification methods are disclosed in several publications and textbooks (Schalkhammer, 2002; Wong, 1991).

According to another aspect of the present invention the substrate can be used for electrophysiological measurements, preferably for patch clamp measurements. Especially apparatus for patch clamp experiments can employ a substrate as disclosed herein.

Preferably the substrate according to the present invention

can be used for a device for simultaneous electrophysiological measurements, preferably for patch clamp measurements, and optical or spectroscopical measurements, if the substrate comprises a transparent electrically insulating material.

Another aspect of the present invention relates to a method for measuring an electrical property of a membranous sample using a substrate as disclosed.

According to a preferred embodiment the membranous sample is selected from the group consisting of cells, cell fragments, cell membranes, liposomes, artificial lipid membranes, lipid layers and mixtures thereof. The membranous sample comprises next to lipids and other cell membrane components particularly transporters and ion channels.

The substrate of the present invention may be preferably used in a method for the simultaneous electrophysiological and optical or spectroscopical measurements of a membranous sample.

Another aspect of the present invention relates to kit for the electrophysiological measurement of a membranous sample comprising a substrate according to the present invention and a device for electrophysiological measurements, preferably a patch clamp device.

The present invention is further illustrated by the following example and figures, yet without being restricted thereto.

Fig.1 shows two examples of substrate - and perforation (a)-structures (sections: 1A and 1B; top view: 1C). The substrate is provided with a plurality of perforations.

Fig.2 shows S-layer coatings. The S-layer adhesion site (a) for the positioning of cells/liposomes/artificial membranes is recrystallized around the aperture (section: 2A, top view: 2B). An inert coating (b) of the residual chip surface can be performed by using another S-layer protein species (section: 2C, top view: 2D). The S-layer adhesion sites (Fig. 2, marked with a) can be structured or non-structured. The S-layer patterns can be performed simple, economical, reproducible and versatile. Structuring can be performed by (i) nano- and micromoulding techniques (Györvary et al., 2003a), (ii) microcontactprinting or (iii) local depletion of previously generated coherent S-layer lattices (microlithography, e.g. ion beam, deep UV) (U.S. Patent: US 6296700 B1). The mould is placed onto the chip surface. The mould can either be prefilled with the distinct S-layer

er solution(s) or the protein solution(s) is (are) injected afterwards into the cavity(ies).

Fig.3 shows possible architectures of S-layer adhesion sites. Areas of tight contact and local specific interactions (= adhesion sites) between the cells/liposomes/artificial membranes may be designed next to each other (e.g. bands, rings, patches etc., hereafter referred to as "zones"). Some examples of possible patterns of S-layers around the chip aperture are illustrated in Fig. 3. The rings, squares/rectangles and bands can be thin (a,f) or broad (b,g). The apertures can be surrounded by a single S-layer zone (a,b,f,g) or by multiple zones (c,d e, h, i, j). The distance between two zones (d,i) as well as the thickness of the different zones (e, j) can be varied. The rings, rectangles or bands can be continuous (a-e, f-j) or non-continuous (k-n). Different zones may be covered with different types of S-layers.

A further structuring embodiment is obtained by time-dependent control of the recrystallization process. Starting from a few randomly distributed nucleation sites, 2-dimensional growth of S-layer domains takes place in all directions, until neighbouring areas meet (Györvary et al., 2003a). Thus, the dimension and the number of crystallites generated per unit area (and hence, the area devoid of S-lattice) can be controlled by the time of incubation (= recrystallization time): Fig. 3; (o) short recrystallization time, (p) longer recrystallization time. When different types (mixtures) of S-layer proteins are used for coating (recrystallization), individual crystallites will assemble next to each other (Sleytr, 1975).

Fig.4 shows the composition of the S-layer lattice (protein type, lattice symmetry, functionality) within the adhesion site (A, B, C, D) and within one zone (E, F, G). Variable architectures (as shown in Fig.3) exhibiting zones of S-layer lattices composed of proteins from either the same (Fig. 4A) or different (Fig. 4B) organisms (native or recombinant; single or variable functionalities) can be used. Different zones of an S-layer adhesion site can be composed of one functionality (Fig. 4C) or variable functionalities (Fig. 4D). One zone can be composed of an S-layer lattice with one single functionality (Fig. 4E) or a mixture of at least two functionalities (same S-layer proteins, but different fused functionalities) (Fig. 4F). One zone can be

composed of S-layer proteins from different organisms exhibiting varying lattice symmetry (pA, pB, pC) and/or functionalities (Fig. 4G).

Fig.5 shows the fabrication of an S-layer adhesion site by using the micromoulding technique (e.g. PDMS mould).

Fig.6 shows additional modification of the substrate surface devoid of S-layer lattice within the adhesion site.

Fig.7 shows S-layer proteins as means for smoothing the sharp corners of the chip perforations.

Fig.8 shows the closing- and opening-behaviour of single gramicidin pores reconstituted in anchored and clamped giant liposomes.

EXAMPLE: The application of 2-dimensional protein crystals for gramicidin ion channel recordings from giant liposomes placed on a planar chip-device

Two-dimensional bacterial surface layer (S-layer) proteins were applied for coating of planar silicon chips and subsequent binding of giant unilamellar vesicles (GUVs) that were functionalized with gramicidin. The GUVs contained a low amount of biotinylated lipids. A genetically engineered S-layer protein that exhibited a fused heterotetrameric streptavidin domain was re-crystallized around the aperture of the microstructured silicon chip forming an S-layer adhesion site. Successful binding via biotin-streptavidin interactions between GUV and the S-layer-streptavidin fusion protein layer (one GUV per aperture) was observed. Gigaseal formation was monitored by resistance measurements. The closing- and opening-behaviour of single gramicidin pores reconstituted in the anchored and clamped giant liposomes was observed. The additional coating of the residual chip surface (around the S-layer streptavidin fusion protein adhesion site) by an inert coating of wildtype SbpA, significantly diminished unspecific adsorption and/or fusion processes of GUVs.

1. MATERIAL AND METHODS

1.1 Chemicals

The phospholipids EggPC and Biotin-cap-DPPE (Avanti Polar Lipids, Alabaster, AL) were dissolved and stored at -20°C in chloroform (Merck, Darmstadt, Germany) at a concentration of 0.1M. As a standard buffer an aqueous solution of 1M KCl was used. The buffer was prepared in MilliQ-water (Millipore, minimum resistance $>18\text{M}\Omega\text{cm}$) and thoroughly degassed and filtered

through a 0.2- μm filter (Sartorius AG, Göttingen, Germany) before use. All chemicals were purchased from Sigma Aldrich (Vienna, Austria).

1.2 Chip (substrate) preparation and treatment

A silicon wafer (NMRC, Cork, Ireland, 1x1 cm², contact angle 64°) was structured by standard lithographic techniques and anisotropic ion etching, resulting in an aperture with a diameter of ~ 1-7 μm . The aperture was placed in the centre of the wafer (one pore/chip). The chips were cleaned by ultrasonic treatment twice for 20 min in 2% Hellmanex (Hellma, Müllheim, Germany) and twice for 20 min in absolute ethanol. After each sonication step, the slides were rinsed thoroughly in MilliQ-water and finally dried in a stream of dry nitrogen. Cleaned chips were stored in clean air until usage.

1.3 Preparation of S-layer protein solutions

Growth, cell wall preparations, and extractions of the S-layer protein SbpA from the gram-positive strain *B. sphaericus* CCM 2177 (Czech Collection of Microorganisms) were performed as described elsewhere (Sleytr et al., 1986). The wildtype SbpA stock solution was diluted with to a final concentration of 0.1 mg/ml. Molecular cloning, gene expression, isolation, refolding and purification procedures of S-layer streptavidin fusion proteins (rSbpA/strp) were described elsewhere (Moll et al., 2002) (Huber et al., submitted). rSbpA/strp proteins recrystallized on hydrophobic silicon as a monolayer with the streptavidin domain exposed to the aqueous solution. An appropriate amount of rSbpA/strp was desintegrated in 4M GHCl for 5 min. The protein was dialysed against MilliQ-water for 2 h. The protein solution was adjusted to a concentration of 1mg/ml with MilliQ-water, centrifuged in an Eppendorf Centrifuge and stored at 4°. For recrystallization rSbpA/strp was diluted with recrystallization buffer (0.5 mM Tris (pH = 9.0) containing 10mM CaCl₂) to a final concentration of 0.1 mg/ml. Recrystallization was performed from 2 to 4h.

1.4 Preparation of giant unilamellar liposomes (GUVs) functionalized with gramicidin

Preparation of giant unilamellar functionalized liposomes was performed by a slightly modified dehydration-rehydration procedure according to Riquelme et al. (1990). In a first step unilamellar proteoliposomes were prepared. Chloroformic solu-

tions of EggPC and Biotin-cap-DPPE (molar ratio 98:2) were added into a glass flask. An ethanolic solution of Gramicidin D, which predominantly (80%) consists of gramicidin A (Sigma, Deisenhofen, Germany) was added (molar lipid/protein-ratio = 200:1). The organic solvent was removed in a rotary evaporator for at least 4h. The dried film was hydrated to a final lipid concentration of 1% (w/v) by adding a standard buffer containing 2% CHAPS (Sigma, Vienna, Austria). The detergent was either removed by dialysis or by addition of Bio Beads SM-2 (Sigma, Vienna, Austria). Glass microscopic slides (ELKA, Germany) were cleaned in Hellmanex and absolute ethanol (ultrasonic treatment), thoroughly rinsed with MilliQ-water and dried in a stream of nitrogen. Small drops (~ 20 µl) or wide stripes (for larger amounts) of the vesicle suspension were deposited onto the glass slides. The suspension was partially dehydrated for 3 h/4°C in a desiccator. Giant liposomes exhibiting a size distribution from 20 µm to 80 µm were obtained by rehydration: an appropriate amount of standard buffer solution was deposited on the top of the dehydrated drop. Size and size distribution of GUVs were examined by transmission electron microscopy (Philips, Eindhoven, The Netherlands) of negatively stained samples. Formvar/carbon coated EM copper grids (Athene old/300 mesh) were deposited on a drop of the sample (5min). Afterwards the sample was fixed with 2.5% glutaraldehyde in 0.1 M sodium cacodylate buffer, pH 7.0 and stained in 2% uranyl acetate.

1.5 Recrystallization of S-layer proteins on the chip

An S-layer adhesion site composed of rSbpA/strp was recrystallized around the micromachined aperture on the chip by a soft-lithographic technique, micromoulding in capillaries, as previously described (Györvary et al., 2003a). For mold formation, mesastructures mold masters in photoresist (AZ 9260; Clariant) on 4-in. silicon wafers were fabricated by photolithographic methods. A master pattern of two horseshoe-like structures was chosen. The horseshoe structures were arranged to each other such that an almost continuous ring with two inlets was formed. Poly(dimethylsiloxane), PDMS (Sylgard 184; DOW corning), is used to fabricate the molds from these masters using the standard procedure (Kumar et al., 1994) (Kim et al., 1995). The mold was carefully placed onto the manufactured silicon chip (microscopical control) such that the aperture was circularly

surrounded by the two horseshoe-like capillaries. Starting from inlets the microchannels (20 μm channel width) were filled with the rSbpA/strp recrystallization solution. After self-assembly and recrystallization of rSbpA/strp proteins the mold was removed under MilliQ-water, leaving the patterned rSbpA/strp arrays behind. Additionally, in some experiments the residual chip surface was coated with wildtype SbpA, which exhibited low un-specific binding surface properties. SbpA formed a monolayer on hydrophobic silicon. The recrystallization quality and integrity of the S-layer was controlled by Atomic Force Microscopy (AFM) analysis. AFM images were recorded in contact mode in liquid with a Nanoscope III AFM (Nanoscope III, Digital Instruments, Santa Barbara, CA). Oxide-shaped silicon nitride tips (Nano-Probes, Digital Instruments) with a nominal spring constant of 0.06 N/m were used. A 100 mM NaCl solution was used for imaging.

1.6 Experimental set-up

The voltage-clamp setup was mounted on a steel core breadboard (CS-11-4, Newport, Darmstadt, Germany). The headstage of the patch clamp amplifier (EPC 9, HEKA Electronic, Lambrecht/Pfalz, Germany) was fixed on a translation stage (M-426, Newport, Darmstadt, Germany) which could be moved vertically (Motorizer 860 A, Newport, Darmstadt, Germany; regulated DC power supply: Kenwood PR 18-1.2, Tokyo, Japan). The headstage of the amplifier, together with measurement chamber were placed within a grounded Faraday cage. The S-layer covered chip was placed horizontally into a home-made teflon-chamber. The chip was sealed into the chamber by a silicone paste (Bayer AG, Leverkusen, Germany) und thus divided the chamber into two compartments. Ag/AgCl redox electrodes were used for voltage-clamp recordings. The reference electrode was placed in the lower, the working electrode in the upper compartment. The lower compartment comprised a vacuum device in order to apply a suction to the clamped liposome. The S-layer covered surface was facing the upper compartment. During this procedure, desiccation of the S-layer was carefully prevented. After placing the chip, the chamber was filled with the standard buffer solution. A small amount of the freshly prepared biotinylated gramicidin-GUV suspension was pipetted onto the S-layer. A single liposome was bound to the rSbpA/strp adhesion zone. After application of a gentle suction, seal formation was monitored by resistance measurements.

1.7 Signal recording and data acquisition

The current response of a given voltage function was measured at 25°C to calculate the electrophysiological parameter of the clamped liposome patch. Conductance was measured via rectangular voltage pulses of up to 200mV across the membrane. All the data handling (voltage stimulation, data acquisition, data storage, data analysis) was performed on a Power Macintosh 7600/120 computer (Apple Computer GmbH, Ismaning, Germany). The setting of the two built-in Bessel filters of the EPC 9 amplifier for the current-monitor signal was 10 and 1.5 kHz, respectively. The data analysis was performed by the Pulse + PulseFit software 7.89/Quadra (HEKA Elektronik, Lambrecht/Pfalz, Germany). Correlations of conductance vs. time and statistical analysis (ANOVA) were performed by using Jandel SIGMA PLOT program for Windows.

2. RESULTS AND DISCUSSION

A prototype of a planar S-layer-based patch-clamp-chip was implemented. GUVs functionalized with biotinylated lipids and gramicidin were chosen as an appropriate system for single channel recordings. The size of GUVs varied from ~20-80 µm, the size distribution was fairly balanced. In some cases, multilamellar vesicles were obtained, which were not used for the experiments. The GUVs were used immediately after preparation or for later experiments divided into aliquots and stored at -80°C until usage.

The recrystallization of rSbpA/strp and wildtype SbpA was monitored by Atomic Force Microscopy. All types of S-layers used in this example recrystallized on silicon chip as monolayered S-layers with the functional (for rSbpA/strp) or the inert surface (for wildtype SbpA) exposed to the aqueous solution. Recrystallization of S-layers on silicon as well as the microstructuring of S-layers by using the micromoulding technique (PDMS moulds) are well-established (Györvary et al., 2003a, 2003b). Diameters of microstructured S-layer adhesion sites ranging from 60-100 µm were obtained. Optionally, the residual chip surface around the rSbpA/strp adhesion site was coated with wildtype SbpA. The outer surface of SbpA is charge-neutral. On chips without a wildtype SbpA layer, adhesion and/or fusion of GUVs to larger lipid aggregates or bilayered structures were observed. A significant decrease in unspecific adsorption and/or fusion processes of GUVs was achieved. Since liposomes only adsorbed slightly and at

a very low amount, they could be washed away easily by gentle rinsing of the upper compartment of the measurement cell. In the majority of experiments a single GUV was bound per S-layer adhesion site. In some experiments either the GUV was not centered and thus no gigaseal was obtained or more smaller GUVs per adhesion site were bound.

Functional studies of single membrane-active peptides and membrane proteins like e.g. ion channels require the reduction of electrical background noise and consequently a high electrical insulation of the surrounding membrane patch, the so-called "giga-seal" ($R > 10^9 \Omega$). The arrangement of the giant liposomes on the perforation is achieved by the linkage of biotinylated lipids within the giant liposomes with the patterned S-layer-streptavidin fusion protein. The tight seal, however, is obtained manually by suction. Stable membranes on the chip with seal resistances of $R > 10^9 \Omega$ were obtained in 17 out of 19 experiments.

The challenge of preparing lipid membranes on silicon chips is the recording of single ion carriers and channels. The general functionality of this concept was demonstrated with gramicidin, a peptide known to self-integrate into one leaflet of a lipid bilayer and to form single channels if dimerization occurs. The closing- and opening-behaviour of single gramicidin pores reconstituted in the anchored and clamped giant liposomes was observed (Fig. 8). As a driving force for ion flux, a voltage typically around 200 mV was applied across the bilayer. Gramicidin is selective for monovalent cations and as KCl solution was used, K^+ currents were recorded. There were clearly five different levels (on horizontal) visible, the lowest representing all channels closed, and each additional level the opening of a channel unit (corresponding to a peptide dimer). Also most channels had the same amplitude of about 8.4 pA, some channels were smaller, i.e., carried less current or transports fewer ions per unit time. This may be due to impurities in Gramicidin D preparation as this preparation is known to contain also a few proportions of other peptides.

The present example shows that stable gigaohm seals over micrometer-sized holes can be obtained. The self-positioning of biotinylated giant liposomes can easily be achieved by the high binding affinity of biotin to the streptavidin heterotetramers which are presented in a well-defined and oriented way by the S-

layer lattice. Thus, no suction is necessary for the positioning of the target molecules, which is always problem as this leads often to clogging of the aperture. If the target molecule is on the right position, suction guarantees the giga-seal formation between the liposome and the aperture. The application potential of this new technique (positioning by the S-layer fusion - lipid anchor molecule interaction and subsequent seal formation by suction of giant liposomes) was clearly demonstrated by measuring the transmembrane current through single ion channels, or in other words by tracing single molecules.

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Claims:

1. A substrate for a device for electrophysiological measurements comprising at least one perforation, characterised in that the substrate is covered at least in part with at least one protein and/or glycoprotein layer.
2. A substrate according to claim 1, characterised in that the at least one protein layer is modified with at least one functionality.
3. A substrate according to claim 2, characterised in that the at least one functionality is selected from the group consisting of proteins, polypeptides, peptides, ligands, chemical groups and mixtures thereof.
4. A substrate according to any one of claims 1 to 3, characterised in that the at least one protein layer is composed of at least one S-layer.
5. A substrate according to any one of claims 1 to 4, characterised in that the substrate comprises an electrically insulating material.
6. A substrate according to claim 5, characterised in that the electrically insulating material is selected from the group consisting of silicon, glass, solid state polymers and mixtures thereof.
7. A substrate according to any one of claims 1 to 6, characterised in that the perforation is 100 nm to 10 μm , preferably 500 nm to 9 μm , more preferably 1 μm to 8 μm , particularly 2 μm to 7 μm wide.
8. A substrate according to any one of claims 1 to 7, characterised in that the substrate is substantially planar.
9. A substrate according to any one of claims 1 to 8, characterised in that the substrate is chemically or physically modified.

10. A substrate according to claim 9, characterised in that the modification is selected from the group consisting of ionizing radiation, atomic radicals, corona treatment, silane groups, chemical groups and mixtures thereof.

11. Use of a substrate according to any one of claims 1 to 10 for a device for electrophysiological measurements, preferably for patch clamp measurements.

12. Use of a substrate according to any one of claims 1 to 10 comprising a transparent and electrically insulating material for a device for simultaneous electrophysiological measurements, preferably for patch clamp measurements, and optical or spectroscopical measurements.

13. A method for the electrophysiological measurement of a membranous sample using a substrate according to any one of claims 1 to 10.

14. A method according to claim 13, characterised in that the membranous sample is selected from the group consisting of cells, cell fragments, cell membranes, liposomes, artificial lipid membranes, lipid layers and mixtures thereof.

15. A method for the simultaneous electrophysiological and optical or spectroscopical measurements of a membranous sample using a substrate according to any one of claims 1 to 10.

16. A kit for the electrophysiological measurement of a membranous sample comprising a substrate according to any one of claims 1 to 10 and a device for electrophysiological measurements, preferably a patch clamp device.

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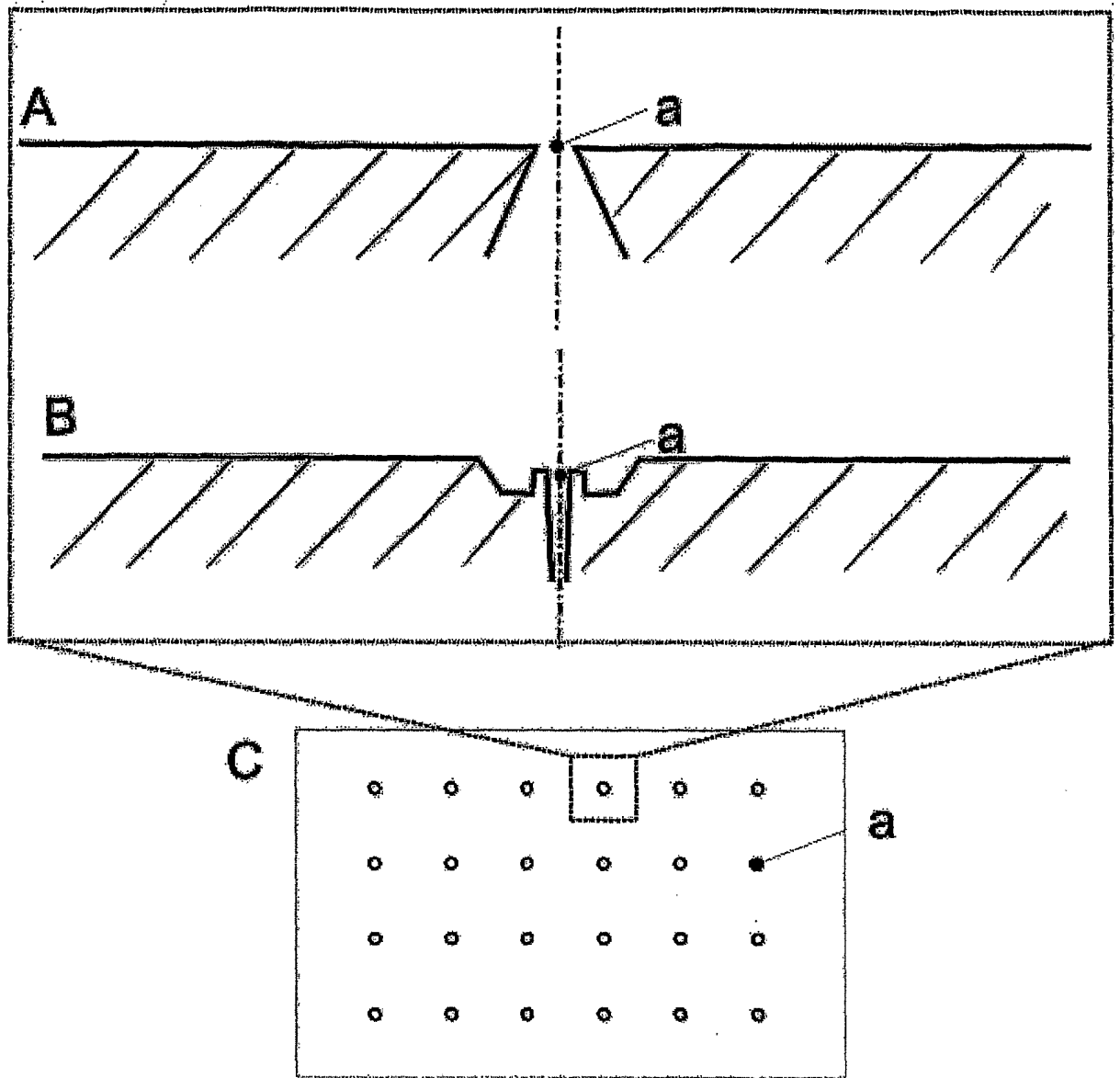


Fig. 1

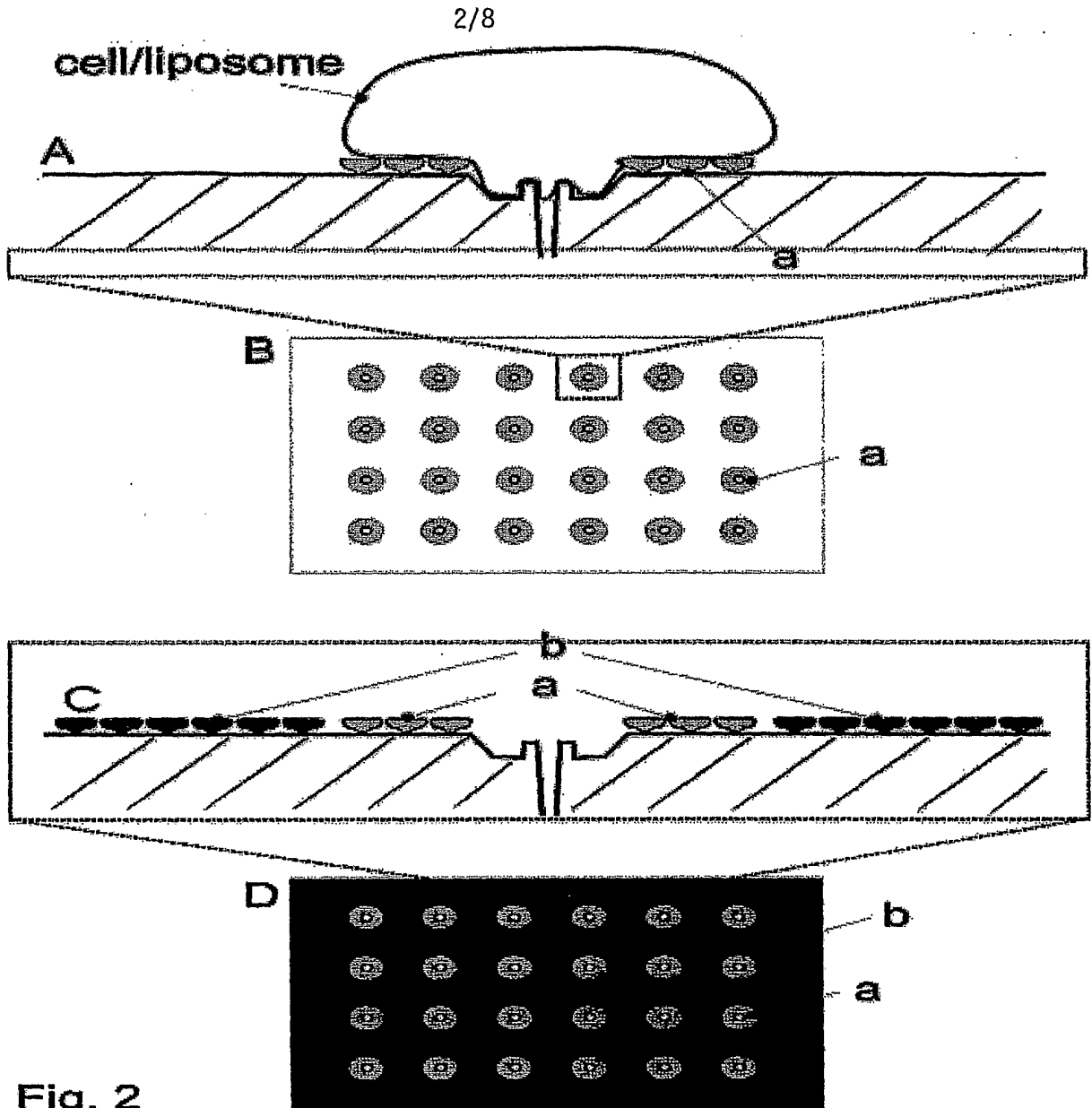


Fig. 2

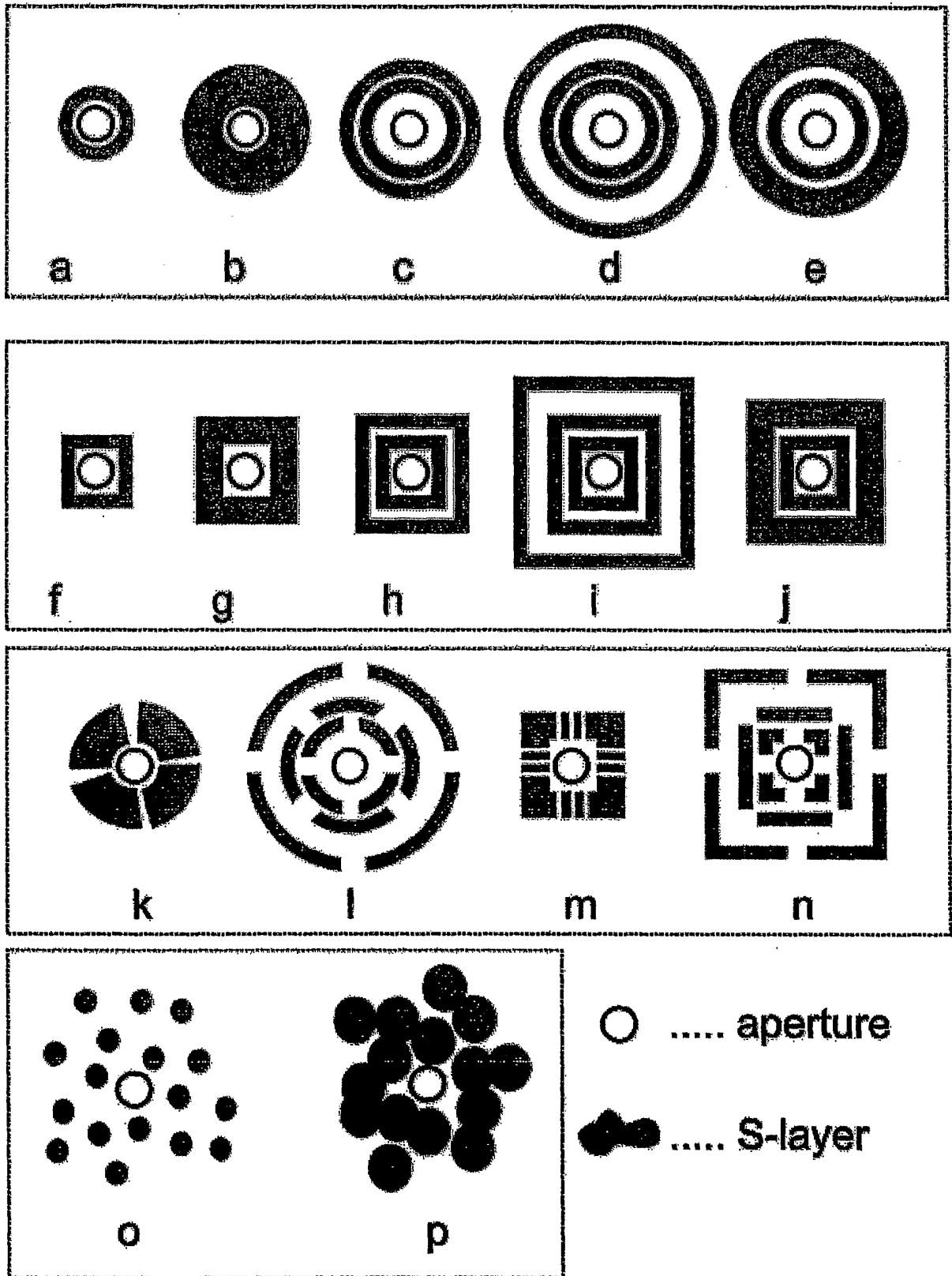


Fig. 3

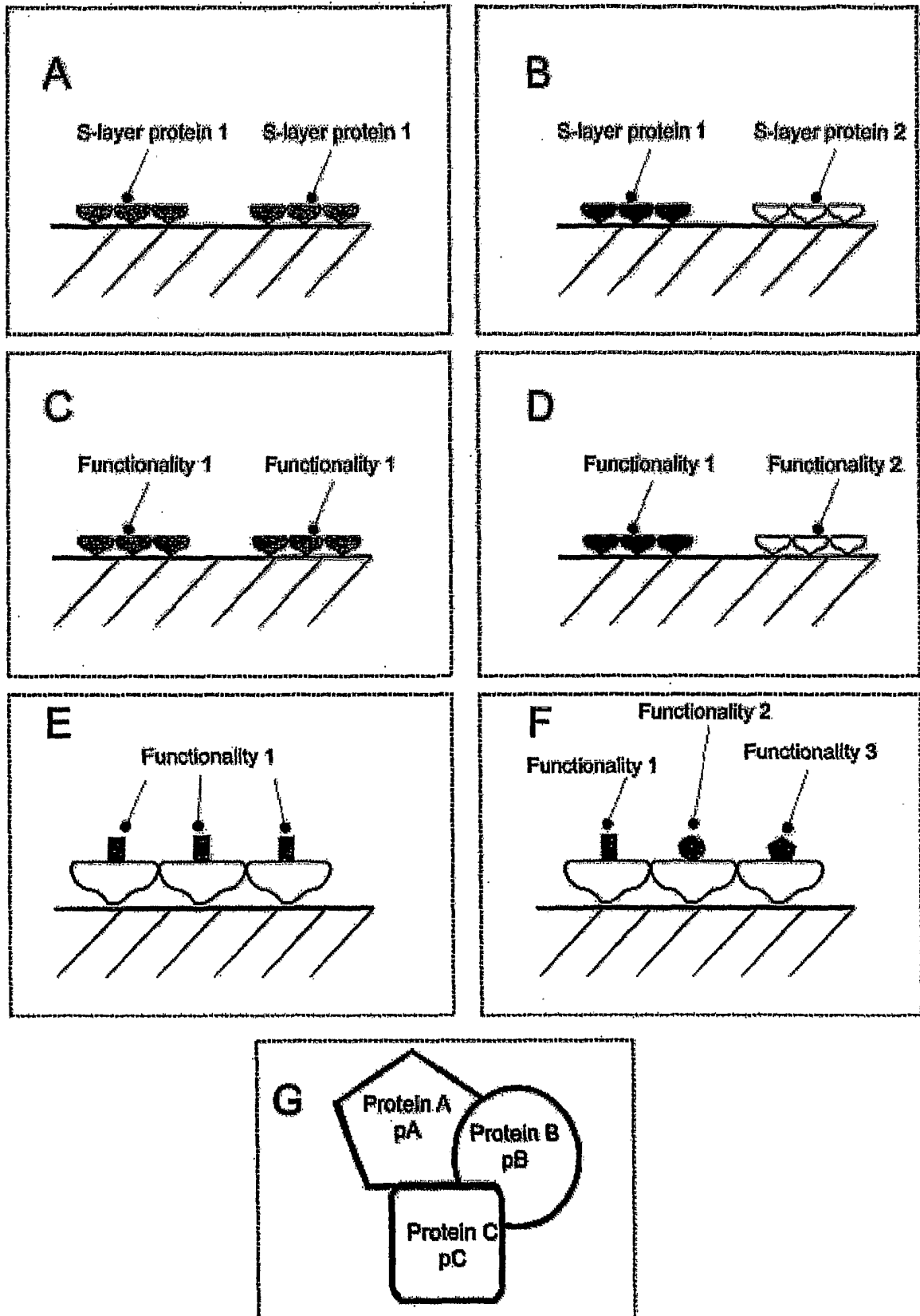


Fig. 4

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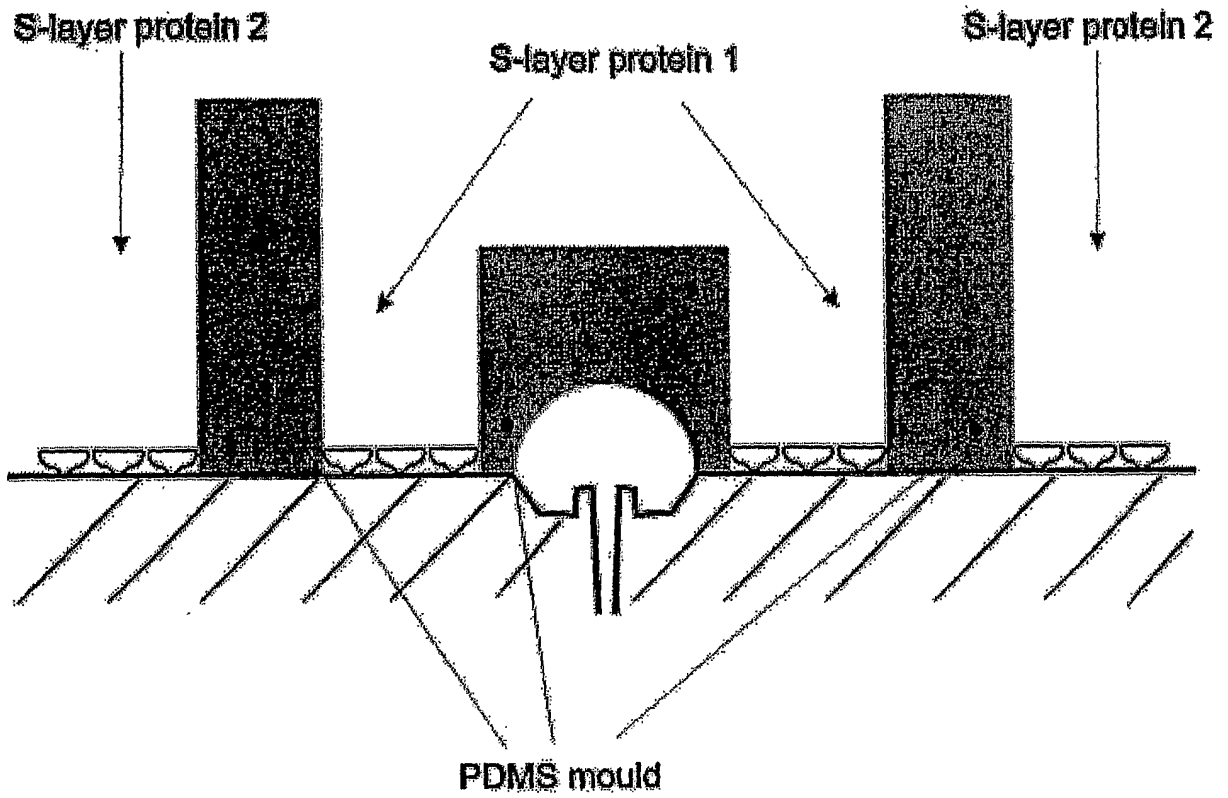


Fig. 5

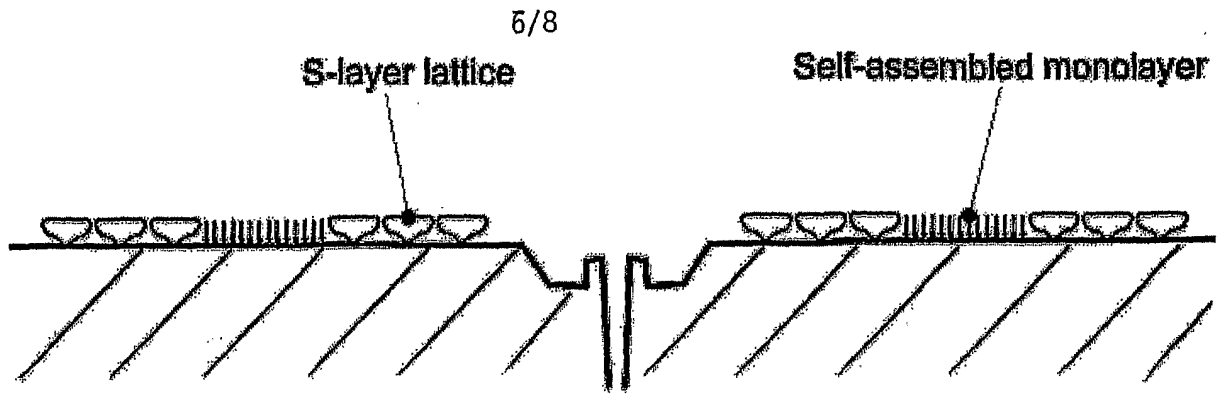


Fig. 6

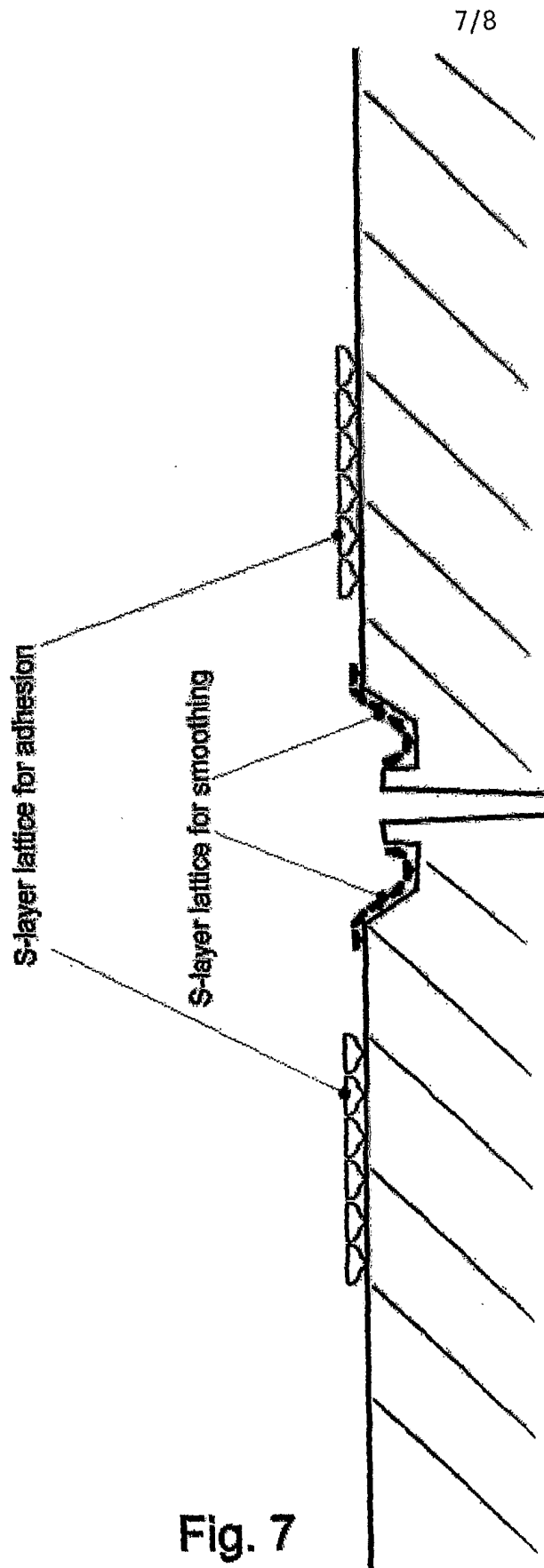


Fig. 7

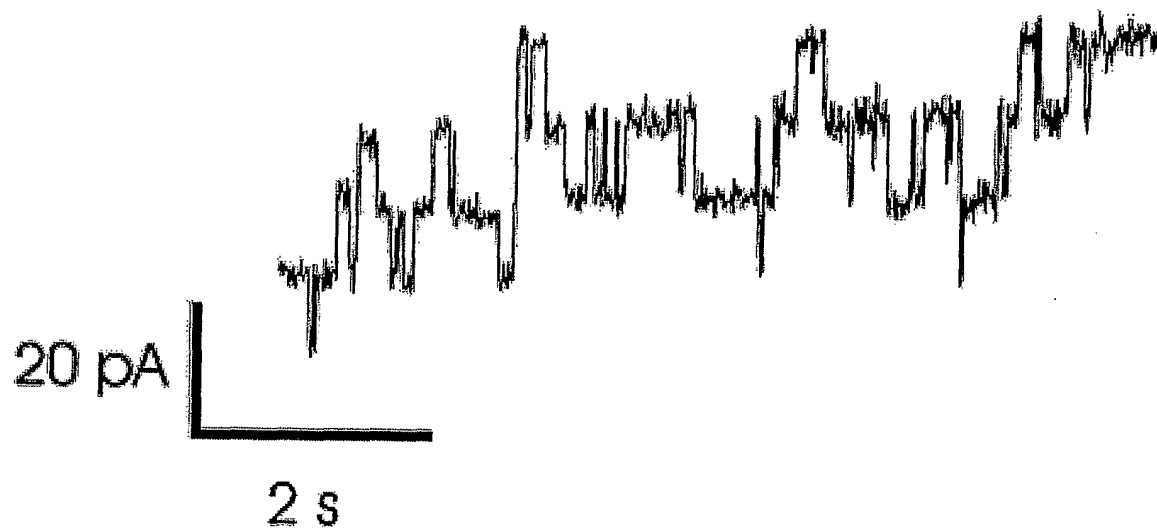


Fig. 8

INTERNATIONAL SEARCH REPORT

International Application No
PCT/EP2004/007878A. CLASSIFICATION OF SUBJECT MATTER
IPC 7 G01N33/487 G01N33/543

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
IPC 7 G01N

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

EPO-Internal, WPI Data, PAJ, EMBASE, BIOSIS, INSPEC, COMPENDEX

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	US 6 315 940 B1 (NISCH WILFRIED ET AL) 13 November 2001 (2001-11-13) abstract figures 1,3 column 5, line 40 - line 45 column 9, line 12 - line 18	1,5-7, 11-16
X	WO 02/04943 A (BRISTOL-MYERS SQUIBB COMPANY) 17 January 2002 (2002-01-17) the whole document in particular page 7, line 14 - page 8, line 11; page 12, lines 4-24; page 13, lines 12-28; Figures 1-3	1-3,5-16
Y	----- -/--	4

 Further documents are listed in the continuation of box C. Patent family members are listed in annex.

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Date of the actual completion of the international search

10 February 2005

Date of mailing of the international search report

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Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2
NL - 2280 HV Rijswijk
Tel. (+31-70) 340-2040, Tx. 31 651 epo nl,
Fax: (+31-70) 340-3016

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INTERNATIONAL SEARCH REPORT

International Application No
PCT/EP2004/007878

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT		
Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	US 2003/146091 A1 (VOGEL HORST ET AL) 7 August 2003 (2003-08-07) the whole document in particular paragraphs 42, 50, 57, 60-67, 82-86	1-3,5-16
Y	-----	4
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Y	-----	4
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Y	----- PUM D ET AL: "The application of bacterial S-layers in molecular nanotechnology" TRENDS IN BIOTECHNOLOGY, ELSEVIER, AMSTERDAM, NL, vol. 17, no. 1, January 1999 (1999-01), pages 8-12, XP004155527 ISSN: 0167-7799 the whole document in particular page 10, col. 1, last paragraph - col. 2, paragraph 2. -----	4

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Information on patent family members

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