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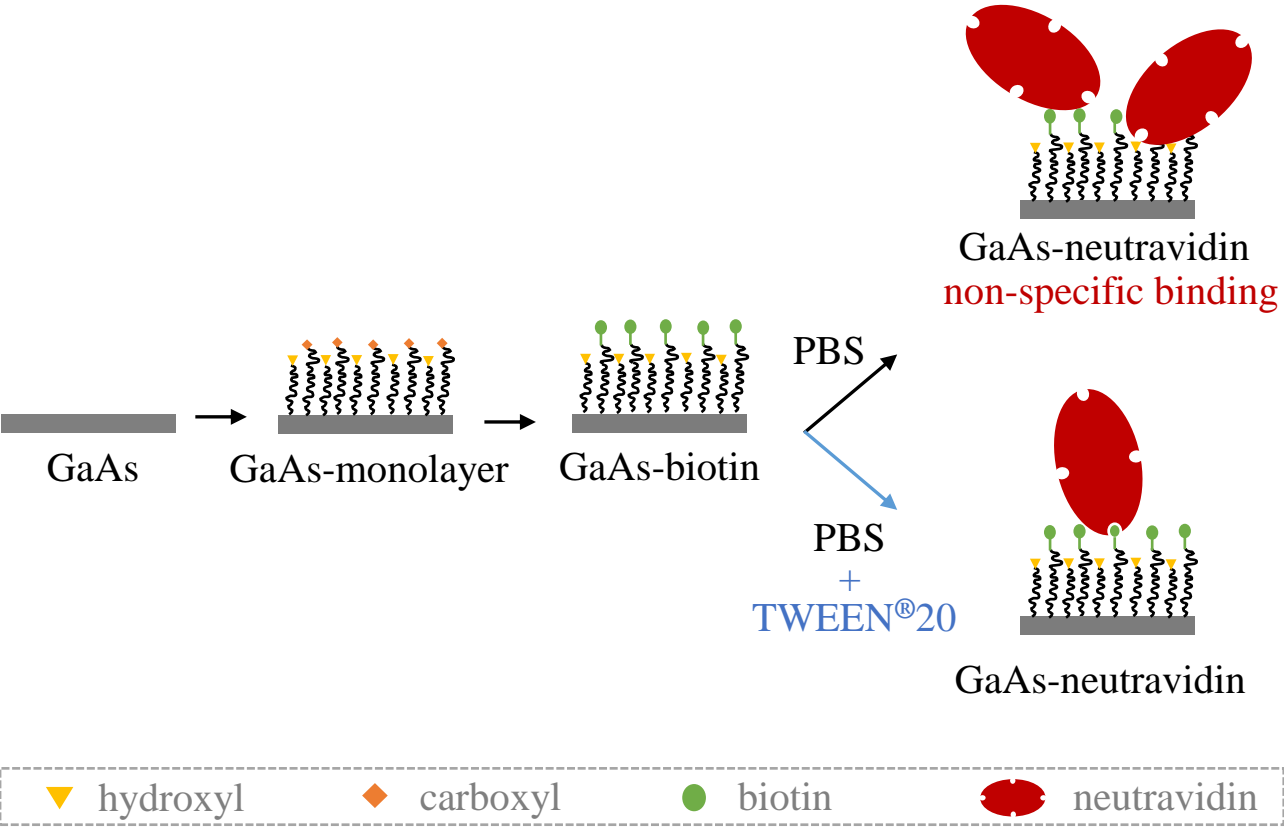
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Graphical Abstract

Biofunctionalisation of gallium arsenide with neutravidin

Bárbara Santos Gomes, David J. Morgan, Wolfgang Langbein, Paola Borri, Francesco Masia



Biofunctionalisation of gallium arsenide with neutravidin

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Abstract

Gallium arsenide (GaAs) is a promising candidate as a platform for optical biosensing devices due to its enabling optoelectronic properties. However, the biofunctionalisation of the GaAs surface has not received much attention compared to gold, carbon and silicon surfaces. Here we report a study presenting a physicochemical surface characterisation of the GaAs surface along the functionalisation with a high-affinity bioconjugation pair widely explored in the life sciences – biotin and neutravidin. Combined X-ray photoelectron spectroscopy (XPS), wettability measurements and spectroscopic ellipsometry were used for a reliable characterisation of the surface functionalisation process. The results suggest that a film with a thickness lower than 10 nm was formed, with a neutravidin to biotin ratio of 1:25 on the GaAs surface. Reduction of non-specific binding of the protein to the surface was achieved by optimising the protein buffer and rinsing steps. This study shows the feasibility of using GaAs as a platform for specific biomolecular recognition, paving the way to a new generation of optoelectronic biosensors.

Keywords: contact angle, x-ray photoelectron spectroscopy, spectroscopic ellipsometry, biotin, self-assembled monolayer, protein non-specific binding, optical biosensor, compound semiconductor, alkanethiolate film, carbodiimide chemistry

1. Introduction

Gallium arsenide (GaAs) is a compound semiconductor material with unique optical and electrical properties, currently used for a wide range of applications, including biosensing devices.[1, 2, 3] In contrast to silicon, whose indirect band gap hinders the integration of light sources in a lab-on-chip design, direct band-gap semiconductors such as GaAs are suited materials for high-speed high-efficiency optoelectronic devices including light sources such as light emitting diodes or lasers. These materials have the potential to provide miniaturised biosensors integrated with CMOS technology, when combined with appropriately engineered bio-components. When developing such biosensors, one crucial step is the functionalisation of the sensing platform with biological recognition elements (so-called bioreceptors).[4, 5] While a variety of systematic studies have been reported for the biofunctionalisation of gold (both flat Au surfaces[6] and Au nanoparticles[7]), silicon[8], and carbon[9], relatively little work has been done exploring the biofunctionalisation of GaAs and the optimisation of such functionalisation strategies, including reduction of non-specific binding. Investigation of passive adsorption of proteins, rather than biospecific interaction of the immobilised ligand with its biological counterpart, is particularly important when functionalising surfaces for biosensing applications.[10]

Nanoscale films with a wide variety of functional groups can be formed by simple self-assembly on surfaces[11], which then allow further functionalisation with bioreceptors.[12] Taking advantage of the highly covalent As–S bond[13], alkanethiolate self-assembled monolayers (SAMs) have previously been formed on GaAs surfaces.[14] These SAMs have been created mostly for GaAs surface electronic and chemical passivation[15, 16, 17] but also for further functionalisation with peptides[18], bacteria[19], virus[20], and proteins.[21] While several studies provided a thorough characterisation of alkanethiolate SAMs on GaAs[22, 23, 24, 25, 26, 27], further functionalisation stages with biomolecules have not been systematically investigated.

This study comprises a detailed physicochemical surface characterisation for GaAs functionalisation with a system widely exploited in the life sciences, biotin/avidin[28, 29, 6, 30, 8]. The popularity of this system stems from (i) the strong and rapid interaction between the biotin and avidin, (ii) the high binding affinity and specificity, (iii) the ability to withstand extreme conditions (in terms of pH, temperature, organic solvents and denaturing reagents) and (iv) the multivalent properties of avidin, with its capacity to bind 4 biotin molecules.

In this study, the GaAs surface is firstly functionalised with oligo(ethyl glycol)-containing (OEG-) alkanethiolate films. Several studies have revealed that the presence of OEG or PEG moieties on the surface can improve the resistance to non-specific binding.[31, 32, 33, 34] However, this ability to resist protein adsorption is not definite and can vary for example according to the molecular conformation of OEG-terminated SAMs formed on those surfaces.[35] SAM formation is followed by covalent attachment of a biotin-based molecule. Subsequent bioconjugation of biotin with its high binding affinity partner – neutravidin – is then promoted. Neutravidin is a deglycosylated version of avidin, which exhibits a lower isoelectronic point and lower nonspecific physical adsorption than avidin, while still retaining the properties mentioned above.

Here we use combined X-ray photoelectron spectroscopy (XPS), ellipsometry and wettability measurements to characterise the GaAs surface at different stages of functionalisation and to investigate the ability of this dual-component SAM to resist non-specific binding of neutravidin. We also investigate how the addition of TWEEN®20 to the buffer where neutravidin is diluted, affects the non-specific binding. TWEEN®20, or polysorbate 20, is a non-ionic detergent that has been used as a blocking agent in various immunoassay methods such as ELISA and western blotting [36]. Incorporated in the washing solutions, it prevents non-specific binding by blocking the vacant binding sites of the microwells after functionalisation with a specific ligand. These molecules with a dual hydrophobic/hydrophilic nature, have also been used to protect and stabilize therapeutic proteins [37]. By binding to the proteins [38, 39, 40, 41, 42] they help preventing protein-protein associations, and by saturating solid surfaces, such as

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the walls of glass vials, they minimise protein adsorption and subsequent conformational changes and aggregation.

2. Materials and Methods

2.1. Materials

GaAs wafers (prime grade, VGF growth, single crystal, orientation (100) \pm 0.1°, undoped, single side polished, 3" diameter, $625 \pm 25 \mu\text{m}$ thickness) were purchased from Wafer Technology. PBS (phosphate-buffered saline 1x, pH 7.4), BupHTM MES (2-(N-Morpholino)ethanesulfonic acid) buffered saline pack, amine-PEG2-biotin and NeutrAvidinTM biotin-binding protein (neutravidin) were purchased from Fisher Scientific. Sulfo-NHS (N-hydroxysulfosuccinimide sodium salt) was purchased from Tokyo Chemical Industry. Unless otherwise specified, chemicals were purchased from Sigma-Aldrich. Solvents of ACS grade were used.

2.2. Methods

2.2.1. Sample preparation

GaAs wafers were cleaved into (5 \times 5) mm² pieces using a diamond scribe. The pieces were cleaned with subsequent ultrasonic baths (5 min each) in acetone, ethanol and isopropanol, respectively, and dried with N₂ flow. Immediately before functionalisation with thiols, the GaAs oxide layer was removed by an acidic bath (2 min in HCl 1 M) followed by rinsing with water and ethanol. Changes in surface roughness after oxide layer removal from GaAs have been previously investigated [43] and it is expected that the surface remains smooth after this short treatment with HCl 1 M. For SAM formation, the samples were incubated in a freshly prepared thiol solution for 24 h, under gentle agitation using an orbital shaker. A mixture of ethanol and water (2:1) with 10% of acetic acid was used as solvent for the two thiols (carboxyl-OEG-thiol, CAS 866889-02-3, and hydroxyl-OEG-thiol, CAS 130727-41-2), which were in solution at 1:1 molar ratio, at a final concentration of 50 μM for each thiol. The use of a mixture of water and ethanol as solvent for thiol solutions has been reported to form superior quality mixed alkanethiolate SAMs, when compared to the use of ethanol only. [25] After SAM formation, the samples were rinsed with the same solvent mixture and dried under N₂ flow. For the attachment of the biotin molecule, via carbodiimide chemistry, the samples were incubated for 30 min in a freshly prepared mixture of N-hydroxysulfosuccinimide (NHS) and N-(3-Dimethylaminopropyl)-N'-ethylcarbodiimide hydrochloride (EDC) in MES 0.1 M, pH 4.7. The final concentration in solution was 50 mM of sulfo-NHS, and 0.2 M of EDC. The samples were thoroughly rinsed with the same buffer (MES 0.1 M, pH 4.7), to remove excess products and by-products. Afterwards, the samples were rinsed with PBS to raise the surface pH and then incubated with biotin solution (50 $\mu\text{g}/\text{mL}$ in PBS) for 2 h. After incubation with biotin, the samples were rinsed with flowing PBS followed by an ultrasonic bath in PBS for 10 min. The unreacted NHS-ester groups were blocked by incubating the samples in alkaline hydroxylamine solution (1 M, pH 8.5), followed by rinsing with flowing PBS and drying under N₂ flow. For neutravidin attachment, the samples were incubated for 2 h in 50 $\mu\text{g}/\text{mL}$ solution of neutravidin diluted in either PBS or PBST (i.e. PBS with 0.05% TWEEN[®]20, pH 6.3), and then rinsed in a sequence of PBST and PBS baths with agitation (1 min in PBST followed by 4 min in PBS, changing the buffer every minute; then repeating the whole procedure 3 times, for a total rinsing time of 20 min) and drying under N₂ flow. Table 1 summarises the samples at different stages of the functionalisation analysed in this work.

This study combines data from three distinct surface analysis techniques: contact angle, spectroscopic ellipsometry and x-ray photoelectron spectroscopy (XPS), detailed below. Triplicate samples were prepared for contact angle measurements. Ellipsometry measurements

ID	Stage	Neutravidin buffer	Washing buffer
#1	Bare GaAs	–	–
#2	Oxidised GaAs	–	–
#3	GaAs-M	–	–
#4	GaAs-M-B	–	–
#5	GaAs-M-B-N	PBS	PBS
#6	GaAs-M-B-N	PBS	PBS/PBST
#7	GaAs-M-B-N	PBST	PBS/PBST
#8	GaAs-M-N	PBS	PBS
#9	GaAs-M-N	PBS	PBS/PBST
#10	GaAs-M-N	PBST	PBS/PBST

Table 1: Overview of the samples investigated. M denotes monolayer, B biotin, and N neutravidin.

were conducted on three different areas of each sample in order to verify homogeneity of the surface. Given the good reproducibility observed for these two techniques, XPS measurements were acquired once.

2.2.2. Contact Angle

Static contact angle measurements were performed by placing a 2 μL water droplet on the sample surface and capturing an image (with a digital camera Nikon Coolpix L120) from the top (Fig. S3). A reference pattern was included in the picture to calibrate the camera pixel and measure the droplet diameter using ImageJ. Finally, the contact angle (θ) was calculated by numerically solving Eq.(1) which expresses the dependence of θ on the wetted radius (r) and the droplet volume (V), with the assumption the shape of the droplet is a segment of a sphere.[44]

$$\frac{3V}{\pi r^3} = \frac{2 - 3 \cos(\theta) + \cos^3(\theta)}{\sin^3(\theta)} \quad (1)$$

The error of the contact angle measurement (σ_θ) was calculated from Eq.(2), which accounts for the error propagation using the functional dependence in Eq.(1), with the error of the volume (σ_V) being taken as the maximum random error of the pipette (0.013 μL) and the error of the radius (σ_r) estimated as 0.05 mm.

$$\sigma_\theta = \frac{1 + 2 \cos(\theta) + \cos^2(\theta)}{\pi r^3} \sqrt{\sigma_V^2 + \left(3V \frac{\sigma_r}{r}\right)^2} \quad (2)$$

2.2.3. Spectroscopic Ellipsometry

A Woollam M200D ellipsometer was used for acquisition of the ellipsometric parameters (Ψ and Δ , in degrees) as a function of the wavelength (spectral range of 200 to 1000 nm). Δ represents the phase shift of the light reflected from the surface polarized in the plane of incidence relative to the one polarized perpendicular to that plane, and $\tan(\Psi)$ represents the ratio of their amplitudes. Measurements were undertaken at 5 angles of incidence: 65°, 68°, 71°, 74° and 77°, with an acquisition time of 1 s and a beam diameter of approx. 4 mm. For thickness calculation, CompleteEASE software (version 5.13) was used and a model consisting of one or more Cauchy transparent layers on GaAs was applied to a wavelength range of 300 to 850 nm, simultaneously to Δ and Ψ data for the five angles of incidence. The fit range has been selected to discard the absorption of the organic layer (for wavelengths shorter than 300 nm) and to avoid multiple reflections arising from the bottom surface once the GaAs becomes transparent at wavelengths longer than 850 nm. The model comprised a GaAs (100) substrate (optical constants uploaded from the software library, shown in Fig. S4) covered by one or more layers (more details on the number of layers used for each sample are shown in SI, Table S1) described

by a 2-parameter Cauchy dispersion model, where the dependence of the refractive index n on the wavelength λ is given by

$$n(\lambda) = A + \frac{B}{\lambda^2}, \quad (3)$$

and the extinction coefficient k is set to 0. An additional term proportional to λ^{-4} can be added to Eq.(3), but it was found that it did not improve the fit. $A = 1.5$ and $B = 0.01 \mu\text{m}^2$ were used as initial guesses, and the fit used the constrain $B \geq 0$. The initial guess value for thickness was 2.8 nm. When more than one layer was added to the model, only the parameters for the top layer were varied by the fit while the properties of the underlying layers were fixed using the results from the previous steps in the functionalisation. For samples where neutravidin was present, the fitting improved significantly (more than 10% improvement in the mean squared error, MSE) if k was described by an Urbach absorption tail

$$k(\lambda) = k_0 e^{\alpha\lambda}, \quad (4)$$

where k_0 and α are fit parameters that determine the shape of the extinction coefficient dispersion. The MSE was calculated by the software and used for evaluation of the goodness of the fitting.

2.2.4. X-ray Photoelectron Spectroscopy (XPS)

XPS measurements were performed using a Thermo Scientific K-alpha⁺ system with a monochromatic Al K α source (1486.6 eV). Charge compensation (at low power, to avoid degradation) was applied to prevent sample charging. An analysis area of $(400 \times 400) \mu\text{m}^2$ and a take-off angle of 90° was used. High resolution spectra were acquired at a pass energy of 40 eV (20 scans, dwell time 0.05 s). For the bare GaAs measurements (# 1), argon clusters (cluster size of 150 atoms at 6 keV, giving an energy of 40 eV per atom) were used to remove oxides from the GaAs surface. CasaXPS software (version 2.3.23rev1.1N) was used for peak fitting using a least-squares method. For each selected region of a scan, linear or 2 parameter Tougaard (so called U2 Tougaard) backgrounds were applied and peak models were created using LA (Lorentzian asymmetric) lineshapes. [45, 46] Elemental relative sensitivity factors (RSF) used for the correction of elemental/component areas were selected from the Scofield library. All the scans were calibrated to the C 1s component at 284.8 eV. For the calculation of the C 1s spectral components, the position (i.e. binding energy) of each component was fixed and the full width at half maximum (FWHM) was constrained to be the same for all the components. The area for the N 1s component was calculated by applying position and FWHM constraints (allowing to move ± 0.2 eV) to the overlapping peaks coming from the substrate. These were assessed from the GaAs-M scan and also verified by a high resolution scan of GaAs acquired just after in situ oxide layer removal with argon clusters (not shown).

3. Results and Discussion

We initiate the functionalisation by creating a dual-component alkanethiolate SAM on GaAs, using the components hydroxyl-OEG-thiol and carboxyl-OEG-thiol (molecular structures shown in Fig. S1). The carboxyl groups are used for the immobilisation of a biotin-NH₂ (see Fig. S1c) molecule via carbodiimide crosslinking (see Fig. S2), while the hydroxyl-OEG-thiol is used as a spacer, allowing control over the density of the immobilised biomolecule, and also providing more rigidity to the SAM through the C₁₁H₂₃ alkyl chain. Following the attachment of biotin to the carboxyl groups, via stable amide bonds, the surface is exposed to neutravidin, with consequent immobilisation of the protein on the surface. A schematic representation of the functionalisation strategy is presented in Fig. 1. Commercially available thiol

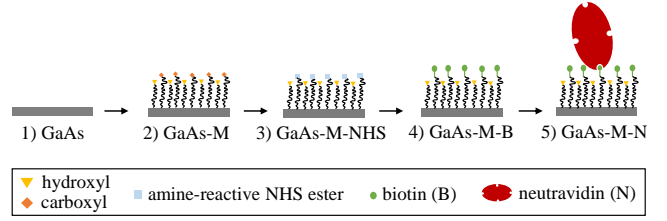


Figure 1: Schematic representation (not to scale) of GaAs functionalisation strategy, showing dual-component alkanethiolate monolayer on GaAs with carboxyl and hydroxyl functional groups (2), intermediate amine-reactive N-hydroxysulfosuccinimide (NHS) ester (3), biotin immobilisation (4), and neutravidin immobilisation (5).

molecules incorporating biotin were not chosen for this study since a thiol with a carboxyl group provides more flexibility for the attachment of any molecules containing amine groups onto the monolayer. The choice of OEG-thiols, rather than polyethylene glycol (PEG)-thiols, is motivated by the low thickness of the functionalised film, suitable for evanescent-wave optical biosensing.

3.1. Wettability

3.1.1. Monolayer formation

Water contact angle measurements were used for investigation of changes in wettability induced by hydrophilic groups (such as carboxyl groups and the aminoacids located on the protein outer surface) immobilised on the GaAs surface. Static water contact angle values for different stages of GaAs functionalisation are shown in Fig. 2. Gallium and arsenic oxides that are formed on the GaAs surface as a result of air exposure [47] were removed by means of an acidic bath, immediately before SAM formation. The oxide-free GaAs (#1, see Table 1) surface is hydrophobic [48], exhibiting a contact angle value higher than 90° . After SAM formation (#3), the measured contact angle was about 67° . The dual-component thiol-based SAM on GaAs was formed by incubation in a solution of equal molar fractions of carboxyl-OEG-thiol and hydroxyl-OEG-thiol. Despite the presence of hydrophilic terminal groups (carboxyl and hydroxyl), the measured contact angle is not below 40° , as commonly reported for carboxyl- and hydroxyl-terminated alkanethiolate SAMs[49, 50]. We attribute this to a contribution from a) the protonation of carboxylate groups[51] and also to b) the ethylene glycol units. Indeed, it has been previously shown, for gold and glass substrates, that OEG-containing SAMs result in a more hydrophobic and flexible surface than the pure alkanethiolate SAMs[52, 53].

3.1.2. Bio-layer formation and detection of non-specific binding

Following biotin attachment (#4), the contact angle value increased slightly (76°) due to less carboxyl groups present on the surface. After protein attachment, the surface became more hydrophilic due to aminoacids being preferentially located on the outer surface of the protein [54, 55]. Varying the protein dilution buffer - PBS (pH 7.4, stage #5) vs. PBST (PBS with 0.05% TWEEN[®]20, pH 6.3, stage #7) - resulted in different contact angle values. Diluting the protein in PBS resulted in a lower contact angle (35°), when compared to PBST (48°), suggesting different amounts of protein present on these surfaces. To confirm if the additional protein coverage observed in stage #5 could be a result of neutravidin non-specific physical adsorption onto the SAM via electrostatic interactions, rather than high-affinity biospecific conjugation of neutravidin to biotin, further investigations were performed and are discussed below.

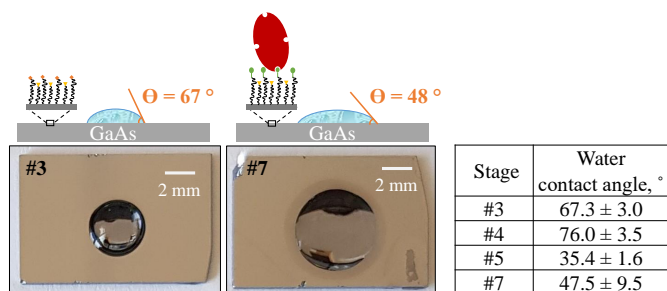


Figure 2: Images (top view) of water droplets on GaAs samples at different stages of functionalisation, with a schematic representation of the drop profile and corresponding contact angle. 10 μ L droplets were used for visualisation purposes. An increase in hydrophilicity was observed after protein attachment. The table presents static water contact angles measured for different functionalisation stages and protein dilution buffer. Sample stages are given in Table 1. The values are the average of measurements on triplicate samples, with the errors representing the standard deviation. The largest measurement error in the contact angle measurement was calculated to be of 0.91°, smaller than the observed standard deviation values.

3.2. Spectroscopic ellipsometry

3.2.1. Film thickness investigation

We investigated the changes in the optical properties of the surface by spectroscopic ellipsometry measurements. This technique measures the change in the polarization of the light upon reflection from the sample surface. Such change is represented by the amplitude ($\tan(\Psi)$) and phase (Δ) of the ratio of the complex reflectance of the p (in the plane of incidence) and s field (perpendicular to p). Film properties such as thickness and refractive index can be calculated from Ψ and Δ using an optical model to fit the data. The spectra of Δ and Ψ for different stages of functionalisation of GaAs are presented in Fig. 3. Bare GaAs (#1) could not be measured as the surface oxidises instantly upon contact with air. Both parameters changed after immobilisation of biotin (#4) and then after incubation with neutravidin (#7), as expected, confirming changes of the surface. As the functionalisation progressed, we observed a decrease in Δ which can be interpreted as an increase in the film thickness, as suggested by fitting the spectra with a multi-layer model. In particular, we observed a decrease in the visibility of the peak around 900 nm, which is a signature of the substrate. Thickness values fitted for these samples are shown in Table 2. Detailed information on the Cauchy model used for each stage, and additional calculated parameters, can be found in SI (Table S1). The thickness value obtained for the GaAs–OH/COOH SAM (#3), 2.3 nm, suggests the formation of a monolayer and is in accordance with a theoretical length of 2.6 nm and 3.3 nm for the fully extended hydroxyl-OEG-thiol and carboxyl-OEG-thiol molecules, respectively. This length was calculated using Chem3D software and is based on the distance measured from the S atom to the farthest C atom in the fully extended molecule. As the functionalisation proceeds, after GaAs–OH/COOH SAM formation, the samples are expected to be composed of more heterogeneous layers, therefore requiring more complex ellipsometric models to fit the data. In addition, for thin films (i.e. with thickness lower than 10 nm) this technique is not as sensitive to changes in the refractive index as it is to variations in the thickness.[56] Nevertheless, an increase in thickness of the films as the functionalisation proceeds is clearly observed.

For optical biosensing exploiting the evanescent wave of light at the GaAs/water interface, it is important that the film thickness is smaller than the decay length of the optical field, which is in the order of 100 nm for light in the near infrared. The measured total thickness of the film does not exceed 10 nm, confirming the suitability of the reported functionalisation strategy.

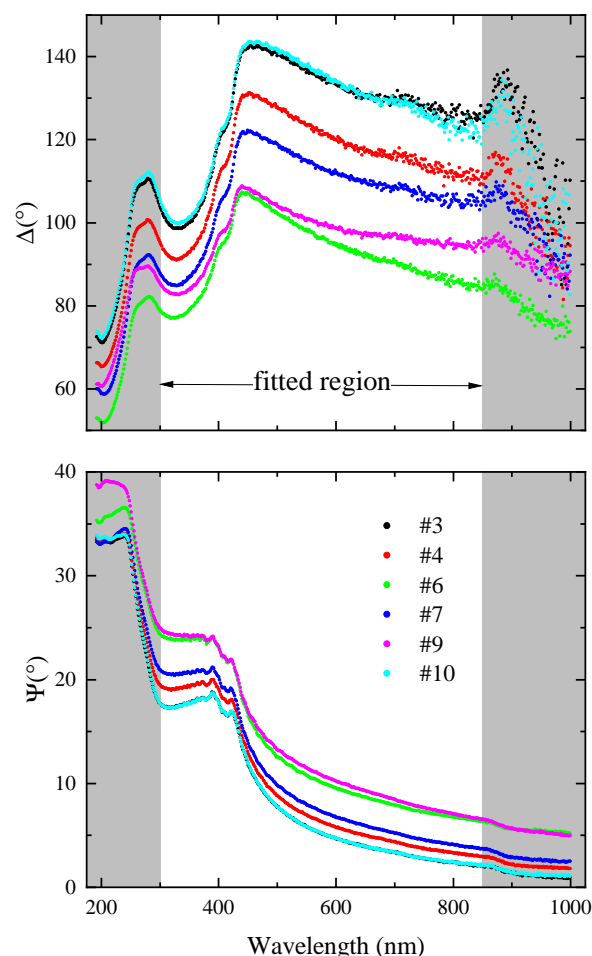


Figure 3: Ellipsometric parameters Ψ and Δ as a function of wavelength, collected at an angle of incidence of 74°, for different stages of GaAs functionalisation, showing changes on GaAs–OH/COOH SAM (#3) after incubation with biotin (#4) and then with neutravidin (#7). Samples stages are given in Table 1. The increased changes observed in #6 with respect to #7 are attributed to non-specific binding, as confirmed by the residual changes measured in #8 compared to the spectra of stages #3 and #10. Spectra collected at additional angles of incidence and corresponding fits are shown in SI Fig. S5 and Fig. S6.

3.2.2. Non-specific binding

A higher thickness (increase of 1.3 nm) was observed for the samples where neutravidin was diluted in PBS (#6), compared to when diluted in PBST (#7), confirming a larger protein coverage. This is in agreement with the wettability results. In order to clarify if the smaller thickness and hydrophilicity detected for the samples with PBST-diluted neutravidin (#7) was due to a reduced attachment of neutravidin to biotin or to a decrease in non-specific binding (i.e. the protein adsorption on random surface locations due to electrostatic interactions), we investigated the adsorption of the protein directly onto the GaAs–OH/COOH SAM. The ellipsometry properties of GaAs–M samples incubated with neutravidin diluted in different buffers were measured. Here, a change in Δ and Ψ is not expected unless the protein binds non-specifically to the SAM, as no biotin is present. The spectra of Δ and Ψ of stage #9 (Fig. 3) showed differences with respect to the GaAs–M (#3), despite the PBS/PBST washes, suggesting a change in thickness caused by non-specific binding of the protein. However, when neutravidin was diluted in PBST (#10) - instead of

Stage	Total thickness, nm
#3	2.31 ± 0.02
#4	4.53 ± 0.17
#6	7.70 ± 0.17
#7	6.41 ± 0.17
#9	7.31 ± 0.65
#10	2.21 ± 0.03

Table 2: Total layer thicknesses at different stages of functionalisation. Sample stages are given in Table 1. Additional information on the models used for the calculation of thickness is given in SI (Table S1). The values are the average of three measurements in different regions of each sample, with the errors representing the standard deviation.

PBS - followed by PBS/PBST washes, the obtained Ψ and Δ curves were similar to the ones of stage #3, indicating an absence of non-specific binding of the protein. These results were further investigated by XPS.

3.3. X-ray Photoelectron Spectroscopy (XPS)

3.3.1. Ratio of neutravidin to biotin on the surface

XPS analysis provided further evidence of functionalisation by providing a chemical signature of the outmost surface layer. Here we present high resolution narrow region scans acquired for two key elements, nitrogen and carbon. These elements are present in all the molecules here used for the functionalisation of GaAs except the thiols, which don't incorporate nitrogen.

High resolution scans for the N 1s region for different stages of functionalisation are shown in Fig. 4. This region is overlapped by Ga $L_{2,3}M_{4,5}M_{4,5}$ (Ga LMM) Auger peaks [57], which complicates quantification of nitrogen species. Nitrogen is absent on the GaAs-M surface (Fig. 4a), as expected, since this element is not present in the thiol molecules (Fig. S1). Biotin attachment is demonstrated by a small shoulder present around 400 eV (Fig. 4b), assigned to C–N bonds [58]. The amount of nitrogen increases after protein attachment (Fig. 4c), as expected, due to the amino acids of the protein. The Ga LMM component overlapping the N 1s signal is very sensitive to oxidation of the samples, which is also reflected in Ga 2p signals (SI Fig. S10). We attribute the higher level of oxidation observed for some of the samples, specifically the samples with neutravidin (Fig. 4c), to a longer exposure to air prior to loading the samples into the spectrometer. Ratios of the areas of the N 1s to Ga $2p_{3/2}$ peaks, calculated from XPS high resolution scans, are shown in Table 3. An increase of N 1s to Ga $2p_{3/2}$ ratio was observed after biotin attachment and then again after incubation with neutravidin, as expected. The observed increase factor of 1.9 between stage #4 and #7 suggests that the ratio of neutravidin to biotin on the surface is approximately 1:25, assuming that each neutravidin molecule has ~ 100 atoms of nitrogen, comparing to 4 nitrogen atoms per biotin-containing molecule. This estimation assumes that the x-ray absorption from the protein layer is negligible and the N signal comes in equal strength from the biotin and the neutravidin layers. This approximation is reasonable considering that the absolute intensity of the Ga peaks measured in stage #4 and #7 are similar (within 20%). The estimated ratio of neutravidin to biotin on the surface is within the expected range, taking into consideration the difference in size for the two molecules. The estimated mass ratio is 246, considering a mass of 244 Da for the biotin molecule and of 60 kDa for neutravidin. If this value is then converted to an area ratio, by simply considering sphere shape for these molecules, the estimated value is of 39. Given the simplicity of this calculation - considering spheres of equal density and similar packing density - a variation of 36 % is plausible.

High resolution scans for the C 1s region for different stages of functionalisation are shown in Fig. 5. This C 1s signal is superim-

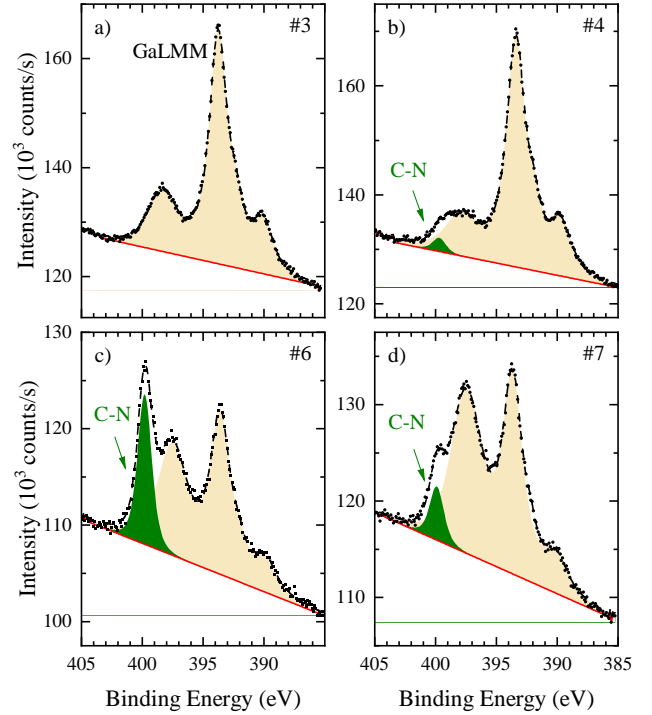


Figure 4: XPS high resolution scans for N 1s region for samples at different point in the functionalisation as indicated by the sample label. The fitted N 1s peak at 399.9 eV is shown in green, while the remaining peaks are assigned to the overlapping Ga LMM (in light brown). The background baseline is shown in red. Nitrogen is absent on the GaAs-M surface (#3), with a small shoulder appearing after biotin attachment (#4); the amount of nitrogen increases after protein attachment, with a smaller nitrogen peak being observed for neutravidin diluted in PBST (#7), comparing with neutravidin diluted in PBS (#6), as a result of decreasing non-specific binding.

posed on the high energy tail of the As LMM peak[59]. As expected, no carbon was detected after in situ argon sputter cleaning applied to the GaAs surface (Fig. S7). For a GaAs sample exposed to the atmosphere, carbon was detected as a result of surface contamination[60] and attributed to C–C and C–H bonds, positioned at 284.8 eV (see Fig. 5a). Formation of the SAM is noted by additional C 1s components (Fig. 5b). Specifically the component around 289 eV is assigned to O–C=O bonds[58], confirming the presence of carboxyl groups, and the component around 286 eV assigned to C–OH and C–O–C bonds, from hydroxyl and ethylene-glycol groups. The concentration of carboxyl groups on the GaAs surface can be tailored by changing the molar fraction of each thiol component in solution (shown in SI, Fig. S8), therefore allowing tailored density of the protein being immobilised on the surface. Scans acquired for the S 2p region (Fig. S9), constitute additional evidence of SAM formation.

The attachment of biotin (Fig. 5c) resulted in a change in the carbon components relative percentages, with an increase in C–C and C–H peaks, and the appearance of a new component at 288 eV, attributed to the carbonyl group (C=O bonds) [58] of the biotin. In addition, an increase of C 1s to Ga $2p_{3/2}$ ratio was observed (Table 3). The increase factor of 1.5 (between stage #3 and #4) suggests that all the carboxyl thiol groups reacted with biotin molecules. This can be explained by the change in the C content of these films as follows. The dual-component SAM formed on the GaAs surface presents equimolar fractions of each thiol (results not shown, from water contact angle measurements of pure SAMs). The number of C atoms is 19 per carboxyl-OEG-thiol, 17 per hydroxyl-OEG-thiol, and 16 per biotin.

Stage	N 1s Ga 2p _{3/2}	C 1s Ga 2p _{3/2}	C 1s components			
			C ₁	C ₂	C ₃	C ₄
#3	(0.003)	1.5	0.75	0.18	(0.01)	0.06
#4	0.10	2.2	0.69	0.28	0.03	(0.004)
#6	0.77	5.3	0.60	0.25	0.15	(0.001)
#7	0.19	2.7	0.63	0.28	0.07	0.02
#8	2.13	10.6	0.56	0.27	0.17	(0.008)
#9	0.68	8.0	0.50	0.39	0.09	0.02
#10	(0.002)	1.8	0.71	0.27	(0.004)	0.01

Table 3: Ratio of N 1s and C 1s peak areas relative to Ga 2p_{3/2}, and C 1s relative component ratios with C₁ detected at 284.8 eV (assigned to C–C, C–H), C₂ detected at 286.2 eV (assigned to C–OH, C–O–C, C–N), C₃ detected at 288.1 eV (assigned to C=O) and C₄ detected at 289.2 eV (assigned to O–C=O). Where no evident N 1s (or C 1s component) peak was present, an upper limit for the ratio was estimated by fitting a peak to the residual; these values are shown in brackets. Additional peak fitting details are shown in Table S2.

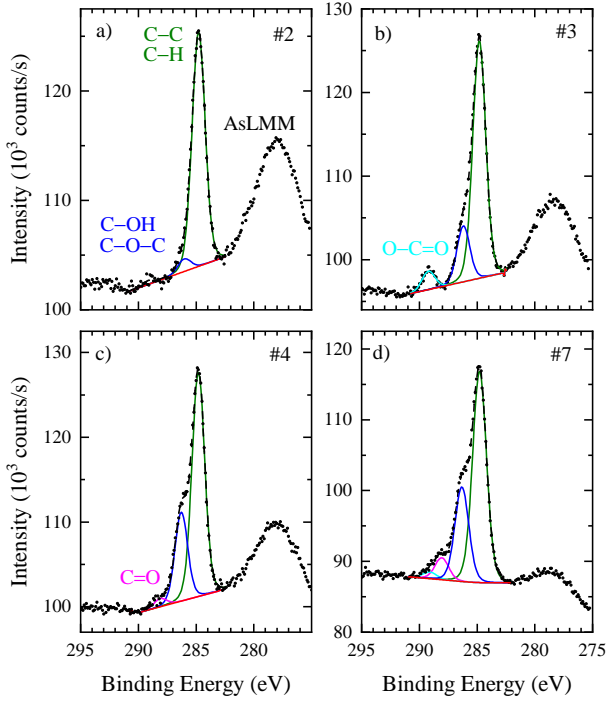


Figure 5: XPS high resolution spectra for C 1s region with fitted components (symbols). This region is overlapped by As LMM auger peaks (not fitted). a) was acquired from a sample of GaAs that had been exposed to air. b-d) are representative scans for different stages of functionalisation of the GaAs samples, as indicated by the sample label. The lines are the fit of the C–C/C–H (green), C–OH/C–O–C (blue), C=O (magenta) and O–C=O (cyan), respectively, with the dashed black line their sum. The background baseline is shown in red. Carbon contamination was detected on GaAs exposed to the atmosphere (#2), with additional carbon components being observed after the SAM formation (#3), confirming the presence of carboxyl groups; the attachment of biotin resulted in a change in the carbon components relative percentages (#4) and this was observed again after neutravidin attachment (#7).

Therefore, an increase factor of $(19 + 17 + 16)/(19 + 17) \approx 1.44$ is expected for C if all the carboxyl groups react with biotin via carbodi-imide chemistry. Further attachment of the protein (Fig. 5 d) resulted, again, in an increase of C 1s to Ga 2p_{3/2} ratio and a change in the relative percentages of carbon components.

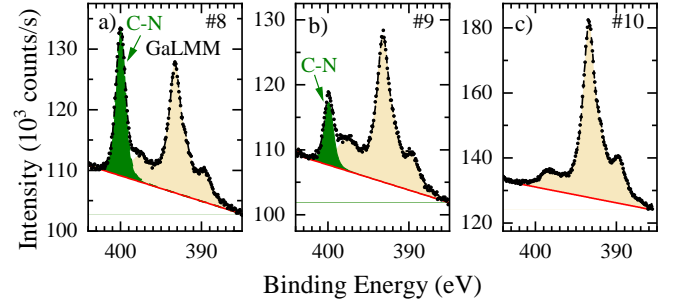


Figure 6: Same as Fig. 4 for samples of GaAs-M incubated with neutravidin as a function of the protein dilution and washing buffers, as indicated by the sample label. The prominent nitrogen peak observed after incubation with neutravidin diluted in PBS (#8) confirms that the protein adsorbed onto the GaAs-M surface; a smaller peak was observed when the samples were rinsed with PBST (#9); in contrast, no nitrogen was detected when neutravidin was diluted in PBST (#10), indicating that in these conditions no protein was adsorbed onto the GaAs-M surface.

3.3.2. Non-specific binding

Fig. 6 shows how the amount of protein adsorbed onto the GaAs-M surface varied as a function of the protein and washing buffers. The prominent nitrogen peak in Fig. 6a confirms that the protein, when diluted in PBS, adsorbed onto the SAM, even in the absence of the specific binding site of the biotin. A smaller but still evident peak was observed when the samples were rinsed with PBST after incubation with the protein diluted in PBS (Fig. 6b). In contrast, when the protein was diluted in PBST and the samples were rinsed with PBS/PBST (Fig. 6c), no nitrogen peak was detected, indicating that no protein adsorbed onto the surface. These results are in agreement with the contact angle and ellipsometry observations, demonstrating the absence of non-specific binding when the protein is diluted in PBST followed by PBS/PBST rinse. This conclusion is also confirmed by the study of the XPS spectra of the C 1s peak (see Fig. S11). We propose that the polysorbate molecules present in PBST buffer bind to certain regions of neutravidin, in a similar way to what has been reported for other proteins [38, 39, 40, 41, 42], and thus reduce the protein surface activity. This prevents the occurrence of electrostatic interactions and hydrogen bonds between neutravidin and the functionalised GaAs surface, while still maintaining affinity to biotin. It is not likely that the polysorbate molecules attach to the functionalised GaAs surface since there was no change in thickness when investigating the attachment of neutravidin diluted in PBST onto the monolayer (#10). It should also be noted that this study has focused on the reduction of non-specific binding of neutravidin and it is expected that the non-specific adsorption will vary in accordance with the physico-chemical properties of the biomolecules being tested. Parameters such as, for example, the pH of the buffer should be adjusted in line with the isoelectric point of the biomolecule of interest.

4. Conclusions

Systematic studies on the biofunctionalisation of GaAs are scarce, comparing to other materials such as gold, carbon and silicon, restraining the opportunity to take advantage of its unique optical and electrical properties for biosensing. We have demonstrated biofunctionalisation of GaAs (100) with a biotin ligand and its biological counterpart, neutravidin, based on combined XPS, contact angle and spectroscopic ellipsometry.

The formation of monolayers was confirmed and the total thickness of the film on GaAs after protein attachment was under 10 nm, validating the suitability of this functionalisation strategy for applications

that require low thickness of the film formed on the substrate, such as optical biosensors that make use of the evanescent wave of light at the GaAs/water interface. The results show that the biotin molecules reacted efficiently with the carboxyl groups on the surface via carbodiimide crosslinking, providing a coverage of neutravidin:biotin with a molecular ratio of 1:25.

Despite the use of OEG-containing SAMs, known to have the ability to resist protein adsorption, [31, 32, 33, 34] we detected significant non-specific adsorption of neutravidin, when using PBS as dilution buffer. However, by incorporating TWEEN®20 - a polysorbate used as a blocking agent in immunoassays methods [36] and as a therapeutic proteins stabilizer [37] - in the protein dilution buffer, non-specific binding was eliminated.

The functionalisation and characterisation strategy herein described can be applied to the immobilisation of other targets on GaAs for biological/biomedical applications. For example, the functionalised film displays neutravidin molecules with up to three binding pockets available for biotin conjugation, and therefore this platform can be used for the immobilisation of biotinylated antibodies or an enzyme or target protein conjugated to biotin [3, 19]. Being composed of a dual-component monolayer, this system provides flexibility for customizing the density of binding sites on the surface.

5. Credit author statement

Barbara Gomes: Writing (original draft), Conceptualisation, Methodology, Investigation; **David Morgan:** Writing (reviewing and editing), Investigation (running XPS samples); **Wolfgang Langbein:** Conceptualisation, Writing (reviewing and editing); **Paola Borri:** Conceptualisation, Writing (reviewing and editing); **Francesco Masia:** Supervising, Conceptualisation, Validation, Writing (reviewing and editing).

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