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Influence of particle size on the SARS-CoV-2 spike protein detection using IgG-capped gold nanoparticles and dynamic light scattering



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ABSTRACT

Due to the unprecedented and ongoing nature of the coronavirus outbreak, the development of rapid immunoassays to detect severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) and its highly contagious variants is an important and challenging task. Here, we report the development of polyclonal antibody-functionalized spherical gold nanoparticle biosensors as well as the influence of the nanoparticle sizes on the immunoassay response to detect the SARS-CoV-2 spike protein by dynamic light scattering. By monitoring the increment in the hydrodynamic diameter (ΔD_H) by dynamic light scattering measurements in the antigen–antibody interaction, SARS-CoV-2 S-protein can be detected in only 5 min. The larger the nanoparticles, the larger ΔD_H in the presence of spike protein. From adsorption isotherm, the calculated binding constant (K_D) was 83 nM and the estimated limit of detection was 13 ng/mL (30 pM). The biosensor was stable up to 90 days at 4 °C. Therefore, the biosensor developed in this work could be potentially applied as a fast and sensible immunoassay to detect SARS-CoV-2 infection in patient samples.

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1. Introduction

The development of fast, low-cost, and easy manipulation diagnostic tests for viruses and related diseases has always been of great interest in science and industry. The SARS-CoV-2 outbreak in late 2019 highlighted, even more, this necessity [1,2]. The recent Omicron variant has infected the population all over the globe with an unprecedented transmission scale and submerging health systems [3]. The most common diagnostic tests for viruses can be divided into three major groups: the amplification of specific

regions of the viral RNA (molecular), rapid antibody tests, and rapid antigen tests [4]. Molecular detection tests, such as PCR (Polymerase Chain Reaction), are the gold standard due to high specificity and sensibility. However, the PCR test involves several processing steps, long analysis time, high cost, and requires highly trained personnel, which can be a bottleneck in terms of scale and population coverage particularly in poor countries. SARS-CoV-2 antibody rapid tests – also referred to as serology – usually detect the antibodies in 17–19 days after symptom onset in the patients, which limits their use for containing pandemic spread [5]. There are available various antigen rapid diagnostic tests (Ag-RDTs), which can detect the SARS-CoV-2 proteins in 15–20 min visually or using small portable devices [6]. These tests are highly specific; however, they are not sensitive as the PCR. Nevertheless, Ag-RDTs can fast

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track and identify COVID-19 disease in individuals. Therefore, the development of cheap Ag-RDTs is of great interest [4,7].

Gold nanoparticles (AuNPs) have been widely explored as transducers in biosensors applied for detection of viruses [8], bacteria [9], metals [10,11], DNA [12], chemical warfare agents [13], contaminants [14], and for the early diagnosis of cancer [15,16] and heart disease [17]. The successful use of AuNPs for sensing applications is mainly related to their facile synthesis and functionalization with various molecules, including the biological ones and their unique optical properties from the localized surface plasmon resonance (LSPR) [18–20]. This phenomenon occurs when the radiation forms a dipole in metallic nanoparticles, with a consequent restoring force to compensate it [21]. The result is a coherent collective oscillation of electrons in a specific resonant frequency [22]. The resonance condition only appears when the frequency of the incident radiation matches the frequency of the oscillating electrons [23]. This frequency depends on the nanoparticle composition, size, and shape, which are the parameters commonly used to control their optical properties. However, the dielectric properties of the environment, the presence of ligands at the surface, and the interparticle approximation generally can have more significant effects on the resonance band than the size [21]. The influence of the surrounding medium or ligands is related to their refractive index, whereas the interparticle distance is important due to the coupling of plasmon resonances, which leads to the formation of the so-called 'hot-spots', broadening the LSPR band and causing electromagnetic field enhancements [24]. The control of all these parameters is key to the application in diagnostics.

When a substance of interest (analyte) causes the aggregation of AuNPs, it induces a color change, which can be visualized with the 'naked' eye to detect its presence. There are also reports of biosensing based on LSPR shift caused by changes in the surrounding refractive index [22,25]. This is only possible because the energy levels of d-d transitions lies in the visible range of the spectrum of AuNPs [26]. The control of color changes is not the only aspect that can be explored from LSPR. This phenomenon also causes an enhancement of the electric fields close to the surface of the particles, making them excellent scatterers [22]. Considering the equation which describes Rayleigh scattering, the intensity of light scattered by small particles ($<\lambda/20$) is proportional to the size of the particles to the sixth power [27]. However, for particles with LSPR, Rayleigh scattering can be many orders of magnitude larger than for non-plasmonic particles. The increase in size results in an increase of the extinction cross-section, and consequently, intensifies the relative contribution of scattering. For AuNPs with sizes up to 80 nm, the optical extinction is dominated by absorption, but scattering takes place when their sizes are larger than that [28–30].

Dynamic light scattering (DLS) also known as photon correlation spectroscopy or quasi-elastic light scattering is a versatile, sensible, and easy operation technique that gives size distributions of colloidal systems [31,32]. The combination of DLS with plasmonic nanoparticles as detection method has receiving attention since the pioneering work of Qun Huo et al. in 2008 on the quantification of prostate specific antigen for cancer diagnosis [33]. They reported AuNPs-based one-step immunoassays capable to detect aggregation occurring in the presence of prostate specific antigen in a 1 ng/mL concentration. Since then, DLS and plasmonic nanoparticles have been used for the detection of many other species, such as other cancer biomarkers [34,35], DNA [36], glucose [37], acetylcholinesterase inhibitors [38], drugs [39,40], metal ions [31,41,42], and particles in gases [43]. Regarding virus detection, DLS-based biosensors have been less explored. The open literature reports successful detection of influenza (H1N1) [8], hepatitis B [44], and SARS-CoV-2 virus [45]. Regarding SARS-CoV-2 protein detection, various biosensors based on the above mentioned LSPR properties

of AuNPs have been recently reported in the open literature, such as colorimetric [46,47], electrochemical [48], electrochemiluminescent [49], opto-microfluidic [50] and so on [51,52]. However, to the best of our knowledge, a different-sized AuNPs study combined with DLS as read-out method for SARS-CoV-2 S-protein detection have not been reported in the open literature.

Herein, the size effect of spherical AuNPs capped with horse polyclonal anti-spike antibody (pAb-AuNPs) on SARS-CoV-2 spike protein (S protein) detection using the DLS technique is reported. The AuNPs were named using their sizes determined using transmission electron microscopy (TEM): **S44** (44 ± 4 nm), **S64** (64 ± 5 nm), **S87** (87 ± 10 nm), **S101** (101 ± 10 nm), and **S130** (130 ± 13 nm). The biosensor response was evaluated by placing the pAb-AuNPs with S protein and reading the average hydrodynamic diameter shift (ΔD_H) caused by the antigen-antibody recognition. The ΔD_H was correlated to the concentration of S protein in the medium and the largest nanoparticle **pAb-S130** provided the best results. Also, we determine the stability, reproducibility, binding constant, and limit of detection (LOD) of the **pAb-S130** biosensor.

2. Experimental procedures

2.1. Materials and methods

Chemicals were purchased from Sigma-Aldrich and used as received: gold (III) chloride hydrate (99.995%), sodium citrate dihydrate (99%), boric acid (99.8%), sodium tetraborate decahydrate, 3,3'-dithiobis (sulfosuccinimidyl propionate) – DTSSP, bovine serum albumin lyophilized powder (BSA). Buffer solutions were filtered with polyethersulfone membrane 0.22 μm pore size (Kasvi) prior to the use. AuNPs, **pAb-AuNPs** were stocked at 4 °C in Falcon or low-binding microcentrifuge tubes. All glassware was washed with aqua regia and deionized water prior to the use. Horse hyperimmune serum was produced by Instituto Vital Brasil (IVB-Brazil) after animal inoculation with the SARS-CoV-2 S protein [53]. The polyclonal antibody (pAb) was purified using affinity chromatography and 1 mg/mL solutions were stocked at –18 °C. The trimeric S protein of SARS-CoV-2 produced in stable recombinant HEK293 cells as described in Alvim et al. was provided by the Cell Culture Engineering Laboratory – LECC from the Universidade Federal do Rio de Janeiro - UFRJ [54].

The samples were centrifuged in low binding microcentrifuge tubes using a Mikro 200R (Hettich) centrifuge in two conditions: (i) 9000 rpm (7690 RCF) at 4 °C for 4 min or (ii) 5000 rpm (2370 RCF) at 10 °C for 4 min. All samples were purified at each synthetic step by two centrifugation/washing cycles (borate buffer pH 8.9, 2 mM). For the DLS measurements, a Zetasizer Nano-ZS90 (Malvern) operating at 90° fitted with a 633 nm 'red' laser was used. The samples were placed in 50 μL or 1 mL polystyrene cuvettes and analyzed at room temperature (18 °C). Each measurement was carried with 10 runs of 10 s in triplicate to calculate the average and standard deviation. The hydrodynamic diameters (D_H) and PDI (polydispersity index) reported in this work are the Z-average obtained from the Malvern Zetasizer 8.01.4906 software. Visible spectra of the dispersions were acquired using a Cary 60 UV-Vis Spectrophotometer (Agilent) at room temperature using 1 mL quartz cuvettes. TEM images were obtained in a Tecnai Spirit microscope operating at 120 kV. For citrate-coated AuNPs samples, one drop of the analyzed dispersions was placed on FCF300-Cu/formvar/carbon film grids and air-dried. AuNPs diameters were determined using ImageJ software (1.53e version) and the histograms of particle-size distributions were plotted for each sample. Attenuated total reflection Fourier-transform infrared (ATR-FTIR) spectra were obtained in the range of 4000–500 cm^{-1} with a spectral resolution of 4 cm^{-1} using a Thermo Nicolet iS50

instrument. For ATR-FTIR analysis, **pAb-S44**, **pAb-S64**, **pAb-S87**, **pAb-S101**, **pAb-S130** biosensors with and without BSA the precursors were concentrated by centrifugation and air-dried.

2.2. Citrate-coated gold nanoparticle synthesis

Citrate-coated AuNPs were synthesized using the seeded-growth method with minor modifications [55]. Briefly, to grow Au seeds, 1 mL of 25 mM HAuCl₄ aqueous solution was added into a 500 mL round-bottom flask containing 150 mL of 2.2 mM Na₃C₆H₅O₇ aqueous solution under stirring. The reaction remained under reflux for 30 min, then cooled to 90 °C, followed by the addition of 2 mL of 60 mM Na₃C₆H₅O₇ and 1 mL of 25 mM HAuCl₄ solution. Subsequent additions of HAuCl₄ (1 mL, 25 mM) solutions were repeated for 2 times in an interval of 30 min, resulting in one growing cycle. After each growing cycle, a dilution process was carried out by extracting 55 mL of the reaction medium, followed by the addition of 53 mL of deionized H₂O. Then, a new growing cycle was initiated by adding the same volume of Na₃C₆H₅O₇ (2 mL) and HAuCl₄ (3 × 1 mL). We conducted this procedure 9 times up to reach the desired particle sizes. Each extracted aliquots were characterized by TEM, visible spectroscopy and DLS. The AuNPs were named using their sizes determined by TEM: **S44** (44 ± 4 nm), **S64** (64 ± 5 nm), **S87** (87 ± 10 nm), **S101** (101 ± 10 nm), and **S130** (130 ± 13 nm). The method used to calculate the nanoparticle concentration is described in the Supplementary Data.

2.3. General procedure for the synthesis of anti-spike antibody-coated AuNPs

Each gold nanoparticle dispersion (300 µL of **S44**, **S64**, **S87**, **S101**, and **S130**) was added to a low-binding microcentrifuge tube followed by the addition of deionized H₂O (800 µL). The tube was centrifuged according to the conditions previously described. Then 1.0 mL of the supernatant was carefully removed, followed by the addition of deionized H₂O (750 µL) and sodium borate buffer (50 µL, 50 mM, pH 8.9). DTSSP solution (20 µL, 10 mM) was added to the tube, which was shaken for 45 min at 18 °C. The dispersion was centrifuged and washed with 2 mM sodium borate buffer (2x). Variable volumes of pAb anti-S protein were added to the tubes and incubated for 2 h at 18 °C. To purify, the dispersions were washed with 2 mM sodium borate buffer (2x). Then, sodium borate buffer containing 0.25% w/v BSA solution, previously filtered in polyethersulfone membrane, was added to the dispersions, yielding the anti-spike pAb-coated AuNPs. The samples were stored at 4 °C.

2.4. General procedure for the pAb-AuNP immunoassay

The **pAb-AuNPs** (10 µL), phosphate buffered saline (PBS) solution (80 µL, 150 mM), and variable volumes of S protein (0.65 mg/mL dispersed in PBS and 300 mM of imidazole) in low binding microcentrifuge tubes were incubated for 30 min at 18 °C in an orbital shaker at 300 rpm. The dispersions were transferred to 50 µL polystyrene disposable cuvettes and analyzed by DLS at 18 °C. Control experiments were performed by measuring the hydrodynamic diameters (D_H) of the dispersions without the spike protein (S protein = S Ptn) in the same conditions and medium. The interactions between all biosensors and S protein (immunoassay) were represented as the hydrodynamic size difference between the D_H of **pAb-AuNP-S-protein** and the D_H of the **pAb-AuNP** control experiment ($\Delta D_H = D_H$ of **pAb-AuNP-S-protein** - D_H of the **pAb-AuNP** control experiment). The LOD was calculated from the adsorption isotherm (Fig. 6a) and determined using the first observed point with ΔD_H value 3 times above the standard deviation of the immunoassay without S protein [56].

2.5. Model employed to fit the data obtained from the immunoassay

The immunoassay can be described as an analytical method in which the quantification of an analyte is realized through its recognition by an antibody. More specifically, it involves the reversible binding through non-covalent interactions of a receptor ($R = \text{IgG}$) to a ligand ($L = \text{S Ptn}$), originating an immune complex (RL). The affinity of the ligand by the receptor is given by the dissociation constant, K_D . This constant can be obtained from the ratio $K_{\text{off}}/K_{\text{on}}$ at the equilibrium, Eq. (1) [57–59].



The binding affinity can be determined using an adsorption isotherm, in which a variable amount of the ligand is added to a fixed concentration of the receptor. The formation of an immune complex can be measured by a diversity of methods and the signal detected for $[RL]$ depends on the ligand concentration in a non-linear manner, usually producing sigmoidal curves. A model commonly used to fit the data is the four-parameter logistic (4-PL), also named the Hill equation, Eq. (2) [60,61].

$$D_H = D_{H,0} + \frac{(D_{H,\text{Max}} - D_{H,0})}{\left[1 + \left(K_D/x\right)^n\right]} \quad (2)$$

In Eq. (2), D_H is the hydrodynamic diameter (nm) for a given value of $x = [\text{S Ptn}]$. $D_{H,0}$ is the hydrodynamic diameter when the concentration of S protein is zero (the minimum asymptote). $D_{H,\text{Max}}$ is the D_H for an infinite concentration of S Ptn, or the maximum asymptote. K_D is the dissociation constant, or the concentration that provides the half-response (sometimes referred as EC_{50}), and n is the slope, or the Hill coefficient of cooperativity. When $n > 0$, the curve is ascendant and indicates a stimulatory effect; and when $n < 0$, the curve is descendent and the effect is inhibitory.

As one-step immunoassays often present the hook effect at high concentration of the analyte, their curves present a bell aspect and are not well-described by a simple logistic curve. To overpass this difficult, we employed an approach that consists the sum of two logistic equations, each one with different n , K_D and $D_{H,\text{Max}}$, Eq. (3) [62].

$$D_H = D_{H,0} + \frac{(D_{H,\text{Max}1} - D_{H,0})}{\left[1 + \left(K_{D1}/x\right)^{n1}\right]} + \frac{(D_{H,\text{Max}2} - D_{H,0})}{\left[1 + \left(x/K_{D2}\right)^{n2}\right]} \quad (3)$$

The immunoassay was fitted employing non-linear regression with the software GraphPad Prism version 9.3.1.471.

3. Results and discussions

3.1. Synthesis of AuNPs and pAb-AuNPs and mAb-AuNPs

To study the influence of gold nanoparticle sizes (AuNPs) in the biosensor response monitored by DLS, we synthesized spherical citrate-coated AuNPs with diameters ranging from 44 to 130 nm using the seeded-growth method with minor adaptations (Fig. 1a and b). With this method, monodispersed nanoparticles of various sizes can be synthesized by the kinetic control of the reaction conditions, such as the amount of Au atoms added to the reaction medium [55]. To synthesize the biosensors, we selected five different sizes of the nanoparticles

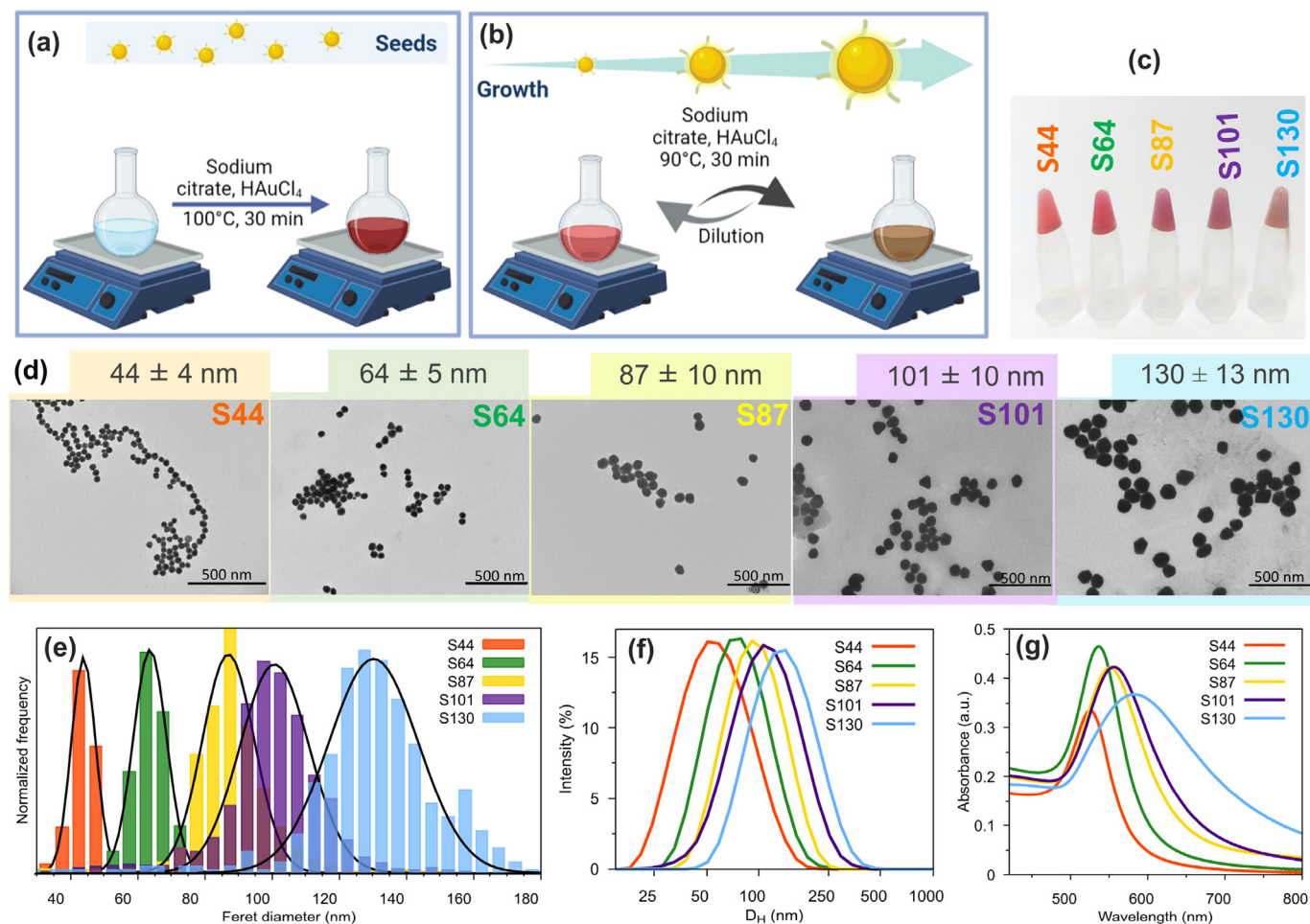


Fig. 1. Synthesis of AuNPs: (a), (b) reaction scheme, (c) photography of the AuNP dispersions as synthesized, (d) TEM images (magnification – 46000 ×, scale bars 500 nm), (e) size distributions and Gaussian fits (TEM), (f) distribution of hydrodynamic diameters (D_H) measured by DLS (dilution in water 1:10), (g) visible spectra (dilution in water 1:10). AuNP, gold nanoparticle; DLS, dynamic light scattering; TEM, transmission electron microscopy.

measured by TEM: **S44** (44 ± 4 nm), **S64** (64 ± 5 nm), **S87** (87 ± 10 nm), **S101** (101 ± 10 nm), and **S130** (130 ± 13 nm). All synthesized AuNPs show monomodal size distributions and spherical shapes, as evidenced by TEM images (Fig. 1d) and DLS analysis (Fig. 1f). The distributions of hydrodynamic diameters (D_H) represented by particle volume and number are also monomodal and without aggregates for all citrate-coated AuNPs (Figs. S3a–e, Supplementary Data). The colors of AuNPs dispersions changed from red to purple and the plasmonic resonance peak broadened and shifted from 525 to 583 nm with their growth (Fig. 1c and g).

The average hydrodynamic diameters (D_H) (Table 1) of all citrate-coated AuNPs obtained from DLS show an interesting result. The nanoparticle diameters of **S44** and **S64** are smaller by TEM than by DLS (D_H), as expected. However, for the larger AuNPs, **S87**, **S101**, and **S130**, the hydrodynamic diameters are smaller than those measured by TEM. Zheng et al. associated this effect to multiple scattering mainly observed for nanoparticles larger than 80 nm in concentrated dispersions [63]. To avoid misinterpretation of the average hydrodynamic diameters of the synthesized AuNPs exceeding 80 nm, we performed serial dilutions of the suspensions and the D_H was measured by DLS (Fig. 2a).

Table 1
Data obtained from citrate-coated AuNPs dispersions and from the biosensors (pAb-AuNPs).

Size	D_H^a AuNP/nm	PDI ^b	λ_{max}^c /nm	[AuNP]/NP /mL	[AuNP]/ μ g/mL	S^d /nm ² /NP	S^d /nm ² /mL	D_H^a DTSSP-AuNP/nm	D_H^a pAb-AuNP/nm (35 μ g/mL)
S44	49 ± 12	0.22	525	21×10^{10}	175	6×10^3	13×10^{14}	50 ± 12	70 ± 18
S64	65 ± 15	0.21	536	8.4×10^{10}	222	13×10^3	11×10^{14}	61 ± 17	85 ± 20
S87	74 ± 18	0.25	552	3.6×10^{10}	241	24×10^3	8.6×10^{14}	91 ± 22	117 ± 28
S101	88 ± 22	0.26	556	2.4×10^{10}	246	32×10^3	7.7×10^{14}	95 ± 25	109 ± 26
S130	121 ± 27	0.19	583	1.1×10^{10}	251	53×10^3	5.8×10^{14}	128 ± 30	146 ± 30

^a Hydrodynamic diameters.

^b Polydispersity index (PDI) of citrate-coated AuNPs.

^c Plasmonic resonance peak.

^d Surface area of AuNPs.

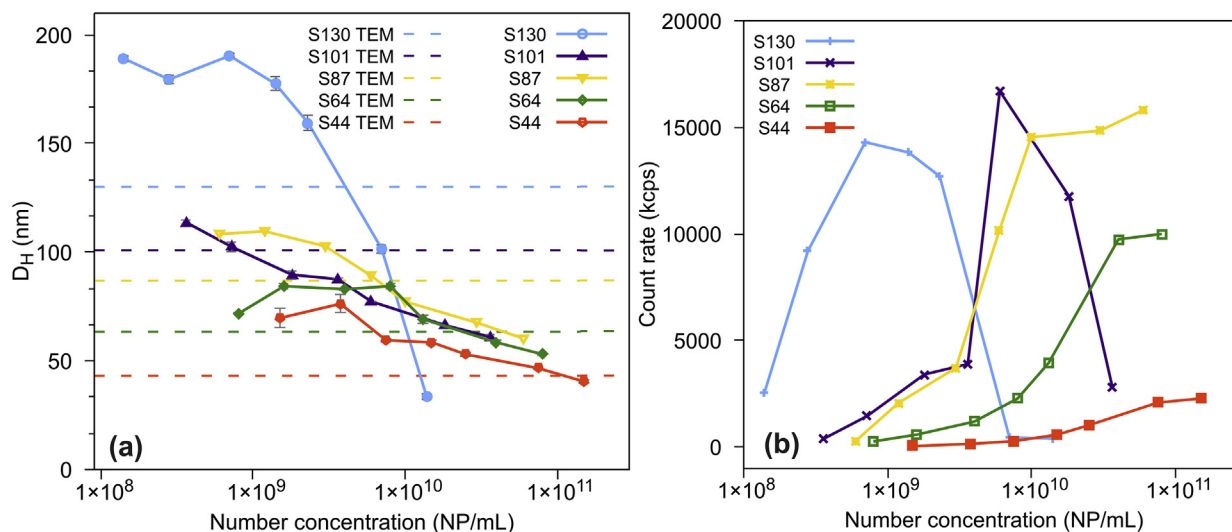


Fig. 2. (a) D_H of the serial dilutions of AuNPs measured by DLS; and (b) scattering (DLS) for the serial dilutions of AuNPs (derived count rate). AuNP, gold nanoparticle; DLS, dynamic light scattering.

The results showed that the D_H was strongly underestimated for more concentrated dispersions of **S101** and **S130**, reaching sizes smaller than those obtained using TEM. Fig. 2b shows that the scattered light intensity detected for **S44–S87** increases with AuNP concentrations. However, for **S101** and **S130**, the scattered light intensity reaches a maximum at 0.6×10^{10} and 0.7×10^9 NP/mL, respectively, and then decreases confirming multiple scattering at higher concentrations. Therefore, for the **S101** and **S130**, we worked at concentrations below 0.6×10^{10} and 0.7×10^9 NP/mL, respectively, to avoid errors in the D_H measurements and, consequently, misinterpreted results due to multiple scattering. For practical applications, in the experimental design in which gold aggregate formation measured using DLS will be used for chemical or biological sensing applications, the optimal concentration of the nanoparticles must be investigated in each case [63]. All DLS dataset obtained from the serial dilutions of the AuNP dispersions are shown in (Figs. S4a–c Supplementary Data).

The concentrations of the AuNPs were calculated in micrograms per mL ($\mu\text{g/mL}$, Au mass) and in nanoparticle per mL (NP/mL) (Table 1). As the particle grows from **S44** to **S130**, the Au concentration ($\mu\text{g/mL}$) in the dispersions increases and the number of particles per unit of volume decreases. As the diameter increases, the surface area of each NP also increases (nm^2/NP). The total surface area should become even larger in the latter cycles of growth. Nonetheless, the number of particles decreases at each dilution cycle because of both volume extraction and HAuCl_4 solution addition. The gold added to the dispersion will recover the particle surface instead of forming new nucleus.

The total surface area per volume unit available to be functionalized ranges from 5.8 to 13×10^{14} nm^2/mL (Table 1). The nanoparticle surface areas (S) do not significantly change with their sizes; therefore, we decided to prepare the biosensors using the same amount of DTSSP ($20 \mu\text{L}$ of a 10 mM aqueous solution) for all AuNPs.

Citrate-coated AuNPs (**S44** to **S130**) reacted with DTSSP. This molecule is a crosslinker broadly used for covalent anchoring of proteins onto gold surfaces [64]. The molecular structure presents a dithiol group in the central moiety, which is cleaved under basic pH and allows the free sulfur atoms strongly bind onto AuNP surface. The hydrodynamic diameters of DTSSP-AuNPs slightly increase compared to the citrate-coated AuNPs (Table 1) because both molecules (citrate and half DTSSP) show similar estimated sizes

(Fig. 3a). The hydrodynamic diameters of the DTSSP-AuNPs were 10% larger (average) than those observed for the citrate-coated AuNPs.

The next step was to bind the polyclonal IgG antibody to DTSSP-AuNPs. The free electron pair of the amino group on IgG can attack the carbonyl group of DTSSP-AuNP forming a stable amide after releasing the N-hydroxysulfosuccinimide leaving group (Fig. S5, Supplementary Data). ATR-FTIR spectra of the **pAb-AuNP** biosensors show the most common absorption bands of proteins, such as Amide I and II from the peptide groups, CH_2 and CH_3 vibrations, and C-NH_2 stretching from Amide III (Fig. S6, Supplementary Data) [69]. Therefore, the ATR-FTIR spectra confirm that the IgG antibody is anchored on AuNP surface.

The optimal concentration for coating polyclonal IgG antibody (pAb) on DTSSP-AuNPs was investigated by ranging the antibody concentration from 5 to $35 \mu\text{g/mL}$ and measuring the D_H (Fig. 3d). The hydrodynamic diameters of all nanoparticles increased as the concentration of the IgG increased, then they stabilized reaching a plateau when the concentration of IgG was around 10 or $20 \mu\text{g/mL}$. The polyclonal IgG antibody concentration employed to prepare the biosensors used in the immunoassay experiments was $35 \mu\text{g/mL}$ because in this concentration, we could ensure that all nanoparticles were recovered. The biosensors (**pAb-S44**, **pAb-S64**, **pAb-S87**, **pAb-S100**, **pAb-S130**) were stored in 0.25% BSA solution prepared in borate buffer (pH 8.9).

3.2. Immunoassay with S protein

To investigate the S protein recognition by the biosensors with different sizes (**pAb-S44**, **pAb-S67**, **pAb-S87**, **pAb-S101**, **pAb-S130**), they were ($10 \mu\text{L}$) incubated with S protein (0.65 mg/mL , $10 \mu\text{L}$) in PBS buffer ($80 \mu\text{L}$) for 30 min , and then their hydrodynamic sizes (D_H) were measured using DLS (Fig. 4a). Control experiments for each biosensor (**pAb-S44**, **pAb-S67**, **pAb-S87**, **pAb-S101**, **pAb-S130**) without S protein in PBS buffer and 300 mM imidazole were carried out and the hydrodynamic sizes were measured to calculate the D_H shift.

Fig. 4a shows that the hydrodynamic sizes (D_H) of the biosensors increase in the presence of 0.65 mg/mL of S protein confirming its recognition by the **pAb-AuNP** systems. The **pAb-S44** biosensor results were discharged because the D_H of its control experiment – the **pAb-S44** dispersion in PBS buffer medium without S protein –

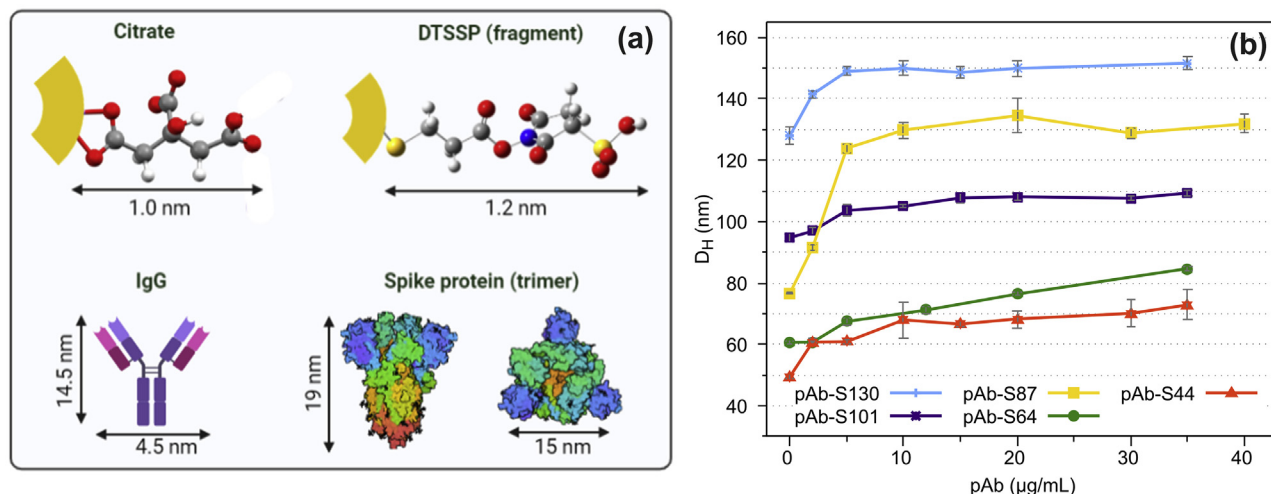


Fig. 3. (a) Estimated sizes to citrate molecule [65], a fragment of the crosslinker (DTSSP) molecule, [66], IgG [67] antibody and spike protein (trimer) [68] and (b) D_H measured by DLS as function of pAb concentrations. DLS, dynamic light scattering; DTSSP, dithiobis(sulfosuccinimidyl propionate); pAb, polyclonal antibody.

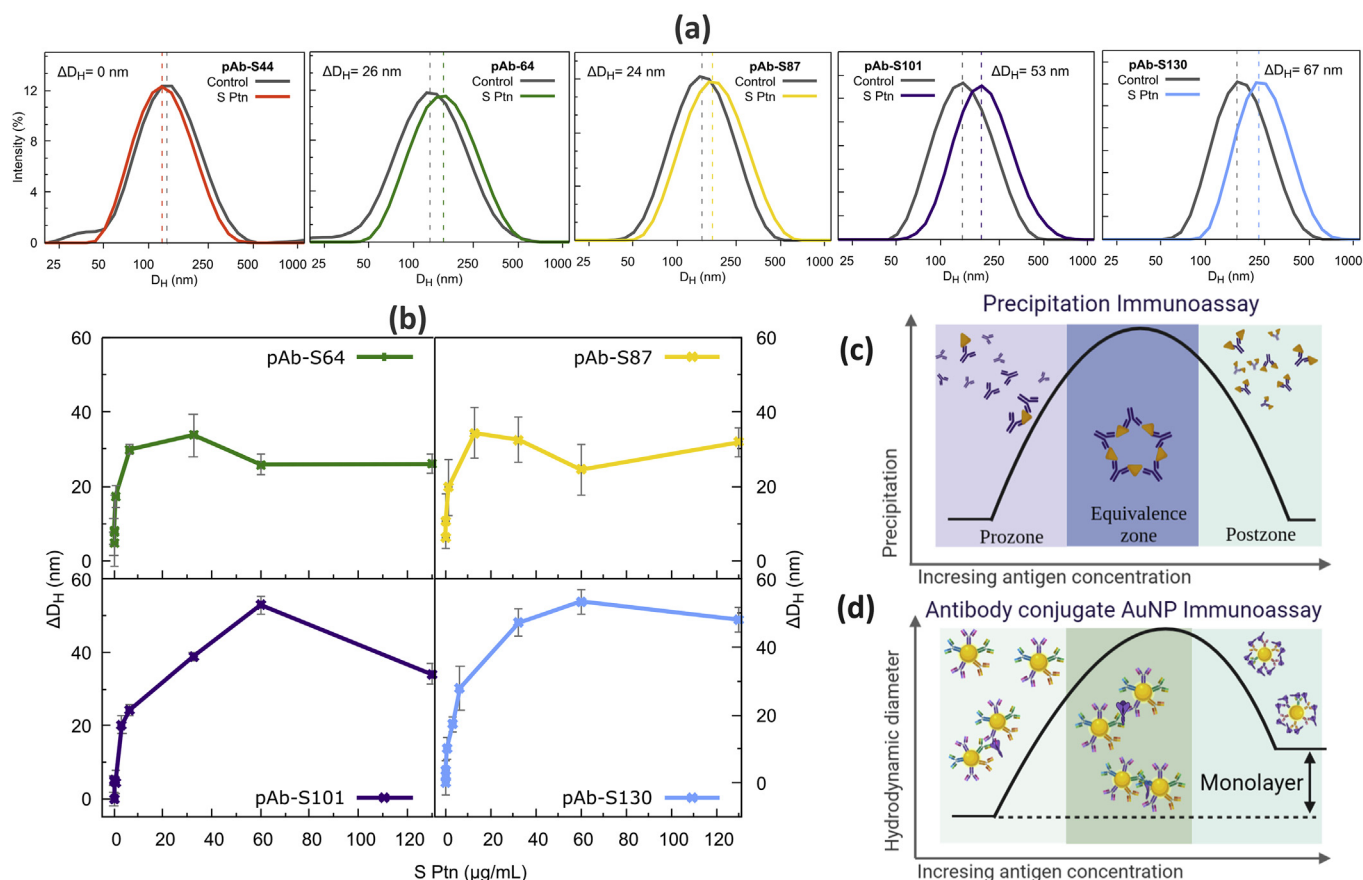


Fig. 4. Immunoassay with S Ptn: (a) ΔD_H measured in the presence of 65 $\mu\text{g/mL}$ S Ptn for all biosensors; (b) S Ptn adsorption isotherm on the pAb-AuNP biosensors; (c) precipitation immunoassay scheme; and (d) scheme showing the D_H variation for antibody conjugate AuNP immunoassay as function of antigen concentration. AuNP, gold nanoparticle; pAb, polyclonal antibody; S Ptn, spike protein.

also increased, resulting in a negligible variation of ΔD_H ($\Delta D_H = D_H$ of pAb-S44-S-protein – D_H of the pAb-S44 control experiment); therefore, the pAb-S44 biosensor is not stable in the investigated medium (PBS buffer and 300 mM imidazole, control experiment). For the larger nanoparticles, the differences between the D_H with S protein and the D_H without it (control), ΔD_H , are bigger than for the

smaller ones. To explain these results, we obtained the adsorption isotherm by measuring the ΔD_H for the biosensors as function of S protein concentration, Fig. 4b.

Fig. 4b shows that pAb-S64 to pAb-S130 biosensors exhibit similar behavior with the increase of S protein concentration, which are an abrupt increase in the ΔD_H with low S protein

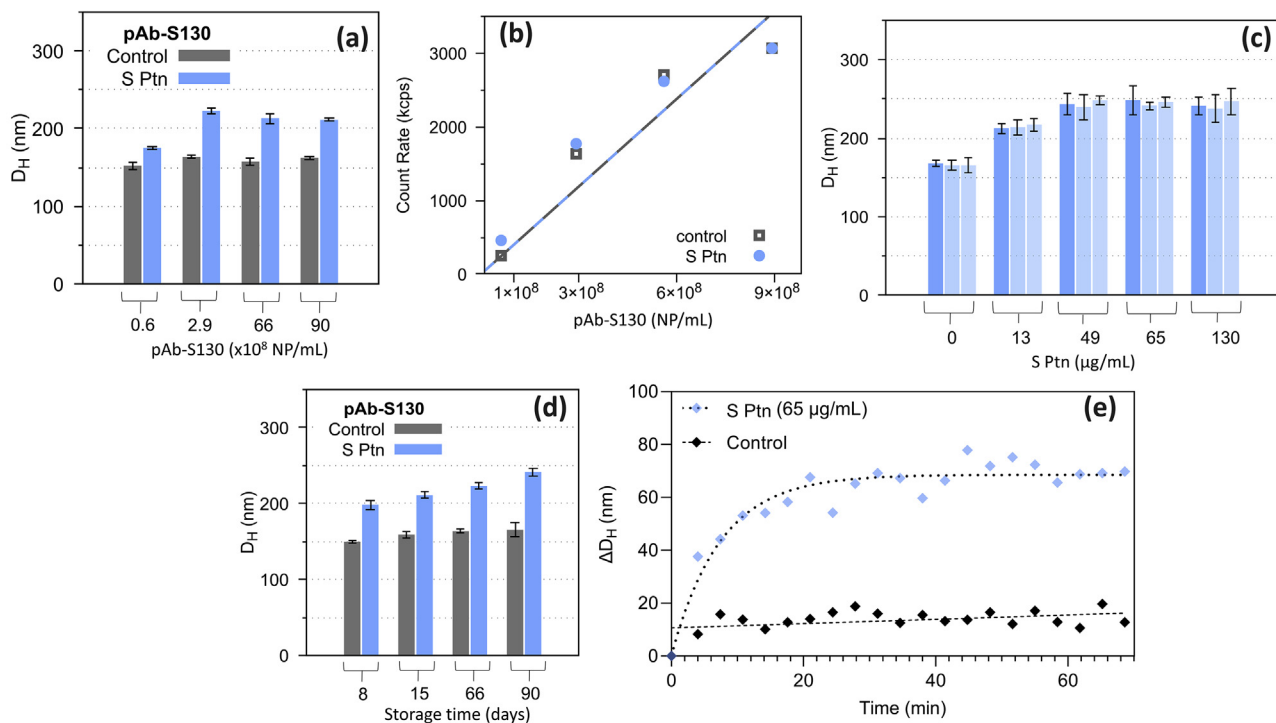


Fig. 5. **pAb-S130** immunoassay: (a) D_H as function of **pAb-S130** concentration without S Ptn (control) and with 65 $\mu\text{g/mL}$ of S Ptn; (b) scattering (count rate) measured for different concentrations of **pAb-S130** without S Ptn (control) and with 65 $\mu\text{g/mL}$ of S Ptn; (c) reproducibility experiment; (d) colloidal stability over the time; (e) ΔD_H measured versus time for **pAb-S130** with 65 $\mu\text{g/mL}$ Ptn S and without S Ptn (control), dashed lines are only eye guides. pAb, polyclonal antibody; S Ptn, spike protein.

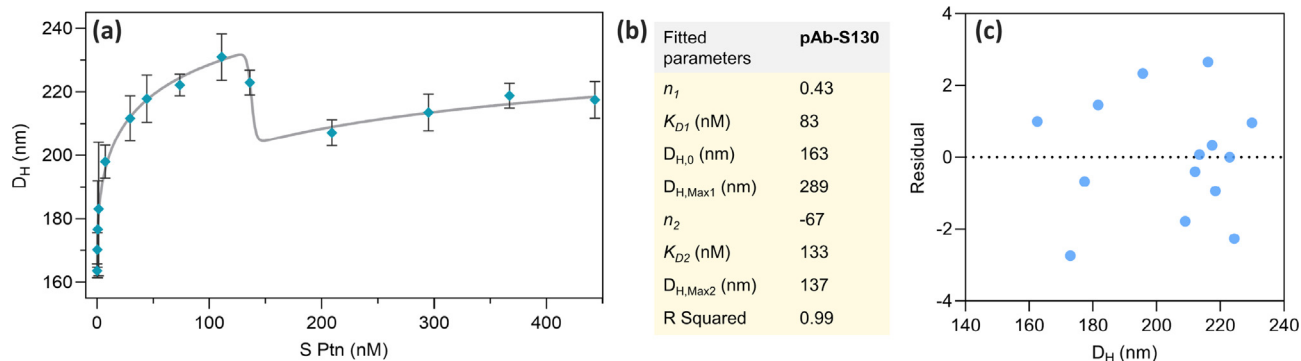


Fig. 6. (a) S Ptn adsorption isotherm on the **pAb-S130** biosensor fitted using Eq. (3); (b) table containing the parameters obtained from Eq. (3); (c) residual plot obtained from the fit of the adsorption isotherm. pAb, polyclonal antibody.

concentrations, followed by a maximum in the ΔD_H , then the values decline and finally stabilize at high S protein concentrations. Such phenomenon is usually called the high-dose hook effect [70].

To illustrate the hook effect, we can make an analogy with the precipitation immunoassay of polyclonal antibodies and proteins, Fig. 4c. In this type of assay, three distinct zones can be defined: (i) the prozone, with the lowest antigen concentration; (ii) the equivalence zone, where the precipitation takes place; and (iii) the postzone, with high antigen concentrations. In the equivalence zone, there is an optimal concentration of both antigen and antibody – which depends on the nature and the specificity of the interaction between them – where crosslinking occurs, and the aggregates are formed. At very high concentrations of the antigen, all interaction sites of the antibodies are occupied, and agglomeration process is inhibited, therefore, precipitation is avoided. When antibodies are anchored on AuNP surface, many factors can affect

the antibody–antigen complex formation. For example, IgG structure has two binding sites in its F_{ab} portion (paratopes) and they sterically influence each other. With the confinement on the nanoparticle surface, the binding sites from neighboring antibodies can influence each other as well. Moreover, the IgG can also adopt different conformations allowing an unpredictable number of paratopes available to interact. Similarly to the precipitation immunoassay, increasing the amount of S protein, an increase in the ΔD_H is noted and then the system reaches a maximum in ΔD_H (Fig. 4b–d). After that, any extra increment of S protein decreases ΔD_H up to reach an approximately constant value. This result can be associated to a postzone-like region in the hook effect, in which there is the formation of S protein monolayer around **pAb-AuNPs** inhibiting their aggregation. In the postzone-like region – where the S protein monolayer is formed around **pAb-AuNPs** – the ΔD_H is in the range of 30–40 nm for the **pAb-S64** to **pAb-S130** biosensors,

these shifts correspond to *ca.* twice the size of the S protein (trimer, see Fig. 3a) [68], Figs. 4b–d.

Furthermore, the ΔD_H was bigger to **pAb-S101** and **pAb-S130** than to the smaller biosensors, Fig. 4a. The agglomeration in the equivalence zone of **pAb-AuNP** immunoassay mainly occurs due to dimer formation, e.g. two **pAb-AuNPs** and one S protein, thus, the larger the nanoparticles, the bigger the dimers [71]. Also, those larger **pAb-AuNP** biosensors require higher concentrations of S protein to reach the postzone region – where the saturation of the binding sites takes place – which accounts for the broader equivalence zone observed in Fig. 4b for **pAb-S101** and **pAb-S130**. Therefore, the **pAb-S101** and **pAb-S130** biosensors show broader antigen quantification regions (dynamic range).

On the light of the mentioned results and considering the higher scattering power of larger AuNPs, we selected the **pAb-S130** biosensor for further investigations. As in the case of **S130** AuNPs, we also analyzed the optimal concentration range of the **pAb-S130** biosensor with S protein (immunoassay) and without it to avoid multiple scattering, which could lead to misinterpretation on the D_H [63]. To find the ideal **pAb-S130** concentration range for the immunoassay, we kept the S protein concentration constant at 65 $\mu\text{g/mL}$ and we changed the concentration of the **pAb-S130** biosensor from 0.6×10^8 to 8.7×10^8 NP/mL and the D_H was measured, Fig. 5a. For the control experiment, D_H was measured without S protein in the same conditions. The D_H of the control experiment did not significantly change as the **pAb-S130** concentration increased, indicating no biosensor aggregation in this concentration range. The count rate of photons for the control experiment and for the immunoassay (Fig. 5b) shows a linear answer from 0.6 to 8.7×10^8 NP/mL evidencing no multiple scattering in this concentration range, therefore, a good working concentration for DLS measurements. For the immunoassay, however, the **pAb-S130** concentration of 0.6×10^8 NP/mL resulted in an almost negligible shift in the D_H with S protein (Fig. 5a). Therefore, considering the interval investigated, the optimal **pAb-S130** biosensor concentration for the immunoassay is in the range from 2.9×10^8 to 8.7×10^8 NP/mL. The **pAb-S130** concentration employed in the following experiments were 2.9×10^8 NP/mL. Fig. S7 (Supplementary Data) shows DLS dataset obtained from the serial dilutions of the **pAb-S130** biosensor.

Fig. 5c shows the D_H reproducibility for **pAb-S130** measured in the presence of different concentrations of S protein. All experiments were performed in triplicates for each concentration of S protein and showed good reproducibility. The colloidal stability of **pAb-S130** biosensor (stored at 4 °C) regarding D_H was monitored over 90 days, Fig. 5d. The results of the control experiment (without S protein) reveal that the D_H of the **pAb-S130** biosensor remains almost the same. Furthermore, upon the addition of S protein (65 $\mu\text{g/mL}$) to the stored biosensor, the D_H increased showing **pAb-S130** remains active for at least 90 days.

We also carried out an experiment to determine the minimum time required to detect D_H variation upon the addition of 65 $\mu\text{g/mL}$ of S protein in the **pAb-S130** biosensor (Fig. 5e). Control experiment without S protein was also conducted. The results show that after 5 min upon the addition of S protein, a shift of 28 nm in the D_H compared to the control experiment (without S-protein) is observed, indicating a fast positive result. After 30 min, the equilibrium was reached and the ΔD_H value was *ca.* 50 nm.

The adsorption isotherm was obtained by measuring the D_H vs. S protein concentration in nM for the **pAb-S130** biosensor, Fig. 6a. To fit the data obtained in this experiment, an adapted Hill equation Eq. (3) was used (see Section 2.5 in the Experimental Procedure) [62]. Considering the ascending part of the curve, we observed a K_D of 83 nM (Table inserted in Fig. 6b). This result is in the same order of magnitude observed for different IgG attached to AuNP

immunoassays reported in the open literature [61]. Also, the K_D obtained in this work is in the same order of magnitude of those reported for llama nanobodies that bind SARS-CoV-2 receptor binding domain in the spike trimer [72]. The Hill coefficient was $n_H = 0.4306$, indicating a stimulatory effect ($n_H > 0$) (Table inserted in Fig. 6b). The maximum D_H value occurs at the concentration of 111 nM of S protein. Above this concentration, the D_H decreases indicating an inhibitory effect. Nearby 200 nM of S protein, a monolayer formation on the biosensor surface takes place, as we illustrated in Fig. 4d. For S protein concentrations higher than 300 nM, a small increase in the D_H values is noted, indicating a multilayer formation tendency. A residual plot (Fig. 6c) shows the selected model satisfactorily fitted the data displayed in Fig. 6a. The LOD was calculated from Fig. 6a using the first observed point with ΔD_H value 3 times above the standard deviation of the immunoassay without S protein [56]. The obtained value for $\Delta D_H = 6.6$ nm was of 13 ng/mL or 30 pM falls under relevant concentration range [73].

4. Conclusions

In this work, we report the development of an one-step detection method for SARS-CoV-2 S protein employing spherical AuNPs capped with horse **pAb** using DLS technique as signal read-out. The AuNPs sizes of the biosensors were analyzed and the larger one, **pAb-S130**, showed the best response. Furthermore, **pAb-S130** demonstrated to be very stable and active for S protein detection up to 90 days and reproducible for different antigen concentrations. Time response experiments showed a positive answer within only 5 min and the maximum ΔD_H within 20 min. The **pAb-S130** biosensor displayed LOD of 13 ng/mL (30 pM). Therefore, due to the affordable, sensitive, user-friendly, robust, and rapid aspect, the proposed method can be considered a potential antigen rapid test.

Credit author statement

The authors of the manuscript gave the following contributions: Carolina B. P. Ligiero, Tamires S. Fernandes, Dayenny L. D'Amato, Francisco V. Gaspar, Paola S. Duarte, Patrícia Bento da Silva, Ricardo B. Azevedo, Glendara Aparecida de Souza Martins, Bráulio S. Archanjo, Camilla D. Buarque, Giovanna Machado, Ana M. Percebom: these authors were responsible for the experimental work up, characterization and S protein detection studies of the different size biosensors, data curation and first draft writing. Marcelo Abrahão Strauch, Juliana G. Fonseca, Leonardo G. R. Meirelles were responsible for the production and biological purification of the horse polyclonal anti-spike antibody used in this work. Célia M. Ronconi: this author was responsible for conceptualization, formal analysis, project administration, supervision and final writing and review.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this article.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.mtchem.2022.100924>.

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