

Surface Plasmon Resonance Immunosensor for Recombinant H1N1 Protein

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Abstract This paper describes about the surface plasmon resonance (SPR) based high affinity detection of recombinant antigen H1N1 (rH1N1HA). The modification of gold SPR chip with 4-MBA and anti-rH1N1HA were in situ characterized by SPR and electrochemical impedance spectroscopy (EIS). By using kinetic evaluation software, K_D and B_{max} values were calculated and found to be 0.278 pM and 84.51 m $^{\circ}$, respectively for the interaction of rH1N1HA with immobilized anti-rH1N1HA. In addition, thermodynamic parameters such as ΔG (−49.78 kJ/mol), ΔH (98.25 kcal/mol), and ΔS (72.59 cal/mol.K) were determined first time for the interaction between anti-rH1N1HA and rH1N1HA, and these values revealed that this interaction is spontaneous, endothermic, and entropy driven one. The kinetics and thermodynamic results of this study revealed that the interaction between the immobilized anti-rH1N1HA with rH1N1HA is more effective than that of interaction of anti-rH1N1HA with the immobilized rH1N1HA. The sensing linearity observed for rH1N1HA is from 0.015 pg to 1.5 ng/ml. The present SPR-based approach may be used in the hospitals equipped with SPR for the early screening of clinical samples in a label-free, real-time manner.

Keywords Surface plasmon resonance · H1N1 virus · Immunosensor · Impedance · rH1N1HA protein

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Introduction

Endemic, epidemic, and pandemic are three words that denote to the spread of infectious diseases among a population, but on different scales. An infection is endemic when it affects a very specific and limited group of individuals at a constant and stable rate. An epidemic occurs when new cases of a certain disease in a given human population, during a given period, substantially exceed the normal rate. Pandemic infections indicate a far higher number of people with a much larger affected region than an epidemic [1–3].

In the case of influenza A virus (H1N1) or swine influenza virus (SIV), it seems suitable to use the term “pandemic” where the number of people with the disease is still small, but the rate of infection is above normal and this disease is present in several continents [4].

In April 2009, a novel influenza A virus (H1N1) was noticed in Mexico and the USA, from which the virus quickly spread across the world [5]. By May 2010, the H1N1 pandemic had caused over 18,000 deaths and had a noteworthy universal economic impact [6]. The 2009 H1N1 influenza pandemic was the first pandemic of the 21st century and a major global public health crisis.

Influenza A viruses are segmented single-stranded RNA viruses from the family *Orthomyxoviridae*. The majority of influenza A viruses infect a broad variety of avian and mammalian species [7]. Influenza viruses are divided into three types: A, B, and C; among these, C is not as important clinically as the A and B types. Virus type A is further subdivided into various subtypes based on the structure and conformation of its two most important envelope glycoproteins: hemagglutinin (HA) and neuraminidase (NA) [8]. HA, the most abundant surface protein consists of three identical monomers with 3 to 9 *N*-linked glycosylation sites. The type of glycosylation and the glycan structure depend on the virus subtype [9]. HA mediates both receptor (glycan) binding and

membrane fusion for cell entry. NA functions as the receptor-demolishing enzyme in virus release [10, 11].

HA is a homo-trimer composed of disulfide-linked polypeptide chains, HA1 and HA2, formed due to cleaving the host's enzymes at the proteolytic site; the former is the main module of the HA antigen. At present, 16 sub-types of HAs and nine subtypes of NAs (N1–N9) have been recognized in avian species (H1–H16) [12–15]. However, the previous four decades have witnessed an increasing number of individual cases of avian influenza virus infections, including H5N1, H7N2, H7N7, and H9N2 sub-types totalling about 50 confirmed transmissions [16–24].

Conventional methods of virus diagnosis including virus isolation, polymerase chain reaction (PCR), and enzyme-linked immunosorbent assay (ELISA) require extensive sample handling, laboratory infrastructure, and long sample-to-result time [25]. Classical methods of influenza virus diagnostics are based on virus isolation, culture, and subtype identification by immunoassays, possibly followed by an *in vivo* experiment to determine pathogenicity [26]. This process requires 3 to 7 days of virus culture in addition to transporting the sample to the laboratory [27]. The foregoing methods necessitate sophisticated bench-top instrumentation, to avoid tedious and time-consuming sample preparation and inclusion of trained personnel. Furthermore, in most of the cases, the amount of sample available for testing is often very small, e.g., blood and cerebrospinal fluid from a neonate. Hence, it is essential to develop a simple, sensitive, and reliable antigen detection technique for the detection of H1N1 virus.

For the direct detection of bimolecular interactions, surface plasmon resonance (SPR) immunosensors have attracted a lot of attention [28] because of its high sensitivity, label-free, and real-time monitoring ability [29]. A novel assay for influenza virus quantification of hemagglutinin (HA) by single-radial immunodiffusion (SRID) using SPR is reported [30]. In addition, SPR technique particularly by combining it with fluorescence spectroscopy was also reported [31, 32].

The use of fluorescence detection schemes in combination with the resonant excitation of surface plasmons was shown to increase the sensitivity for bioanalyte monitoring considerably. The applicability of a new technique, microwave-accelerated surface plasmon-coupled luminescence (MA-SPCL) for fast and sensitive bioassays in buffer, serum, and whole blood using quantum dots as luminescence reporters is also demonstrated [33]. Moreover, SPR sensing of anti-rH1N1HA was reported by our team by immobilizing rH1N1HA on the SPR chip [34]. In continuation to our earlier studies on the development of SPR detection methodologies for BWAs [35–38] in this present work, SPR-based methodology is developed for the detection of rH1N1HA in phosphate buffered saline (PBS) medium using 4-mercaptobenzoic acid (4-MBA) modified gold chip immobilized with anti-rH1N1HA. To the best of our knowledge, so far, no SPR

report is available for the sensing of rH1N1HA with its immobilized anti-rH1N1HA on 4-MBA modified gold SPR chip. This modified gold chip was characterized by SPR and electrochemical impedance spectroscopy (EIS). Parameters affecting the response of SPR were optimized and finally, equilibrium constant (K_D) and maximum binding capacity of analyte (B_{max}) were calculated, and also for the first time, thermodynamic parameters such as change in Gibb's free energy (ΔG), change in enthalpy (ΔH), and change in entropy (ΔS) involved in the interaction between immobilized anti-rH1N1HA and rH1N1HA were also deduced in this study.

Materials and Methods

Chemicals and Reagents

N-(3-dimethylaminopropyl)-*N*-ethylcarbodiimide hydrochloride (EDC), *N*-hydroxysuccinimide (NHS), phosphate buffered saline (PBS), sodium acetate, ethanolamine, and hydrochloric acid (HCl) were of Fluka grade and obtained from Sigma-Aldrich, Bangalore, Karnataka, India. Glacial acetic acid, glycine, sodium hydroxide (NaOH), and methanol (MeOH) were supplied by Sigma-Aldrich, Bangalore, Karnataka, India. 4-MBA Aldrich purchased from Sigma-Aldrich, Bangalore, Karnataka, India was used for the modification of the SPR gold chip (XanTec Bioanalytics GmbH, Metrowingerplatz, Germany). All chemicals and reagents used were of analytical grade, and purification was performed wherever necessary before use.

Moreover, rH1N1HA antigen (M.Wt. 75 kDa) was prepared by the trained biologists as reported earlier [34] in our establishment in addition to the raising of anti-rH1N1HA (M.Wt. 160 kDa). The concentration of rH1N1HA antigen (15 mg/ml) and anti-rH1N1HA (5 mg/ml) were used in this study.

Different buffer solutions were used in this study depending on pH [acetate buffer (pH 4.0–5.5), PBS buffer (pH 6.0–7.5), and glycine–NaOH buffer (pH 8.0–9.0)] for the optimization of pH. All solutions were prepared using water from a Milli-Q system (Millipore India, Bangalore, Karnataka, India) throughout the experiment.

Instruments

The biomolecular interactions were explored using a two channel cuvette-based electrochemical surface plasmon resonance system (Autolab ESPRIT, Ecochemie B.V., The Netherlands) and diode laser is a source with a fixed wavelength of 670 nm in conjunction with a scanning mirror to modulate the plane polarized light beam on the SPR substrate. The outcome of the SPR measurement was automatically monitored using a PC with data acquisition software version 4.3.1. All kinetic

data were obtained using kinetic evaluation software version 5.0 (Ecochemie B.V.). Bare gold chip was modified with 4-MBA by using spin coater (Autolab ESPRIT, Ecochemie B.V., The Netherlands). The pH of the buffers was measured with a EUTECH instruments pH metre (pH 1500, Ayer Rajah Crescent, Singapore). All experiments were carried out at 25 °C unless otherwise specified, and the temperature of cuvette was controlled by a Julabo HE-4 (Julabo Labortechnik GmbH, Seelbach, Germany) water bath.

Preparation of 4-MBA Modified Gold Chip

Bare gold SPR chip was modified with 0.01 M methanolic solution of 4-MBA using a spin coater. Firstly 75 µl of 4-MBA solution was dispensed on bare gold chip and spread at 100 rpm. After this, spin speed was increased up to 2,500 rpm and kept for 5 min to spread the 4-MBA homogeneously on the gold SPR chip.

Immobilization of Anti-rH1N1HA on 4-MBA Modified Gold SPR Chip

Prior to the immobilization of anti-rH1N1HA on 4-MBA modified gold SPR chip, 50 µl of PBS buffer (pH 7.5) was passed every 120 s interval for 60 s in order to get a stable baseline in both the channel. 4-MBA modified gold SPR chip was chemically activated by the injection of 75 µl of a 1:1 mixture of 400 mM EDC and 100 mM NHS. Subsequently, 75 µl of anti-rH1N1HA (1:100 dilution in 10 mM PBS) was injected in channel 1 for 1,800 s to get an effective immobilization of anti-rH1N1HA on the activated 4-MBA modified gold SPR chip surface. Following anti-rH1N1HA immobilization, the remaining active sites were blocked with 75 µl of 1,000 mM ethanolamine. Afterwards, 10 mM HCl was injected to achieve regeneration of immobilized sensor surface. For negative control measurements, the modified gold surface was activated with EDC/NHS and then quenched with ethanolamine (channel 2) as mentioned above and was used as blank control surface. This anti-rH1N1HA that was immobilized on the modified gold SPR chip was interacted with different concentration of rH1N1HA, in order to see and also deduce kinetic and thermodynamic parameters.

Biosensing Protocol

Binding studies was performed using different dilutions of rH1N1HA in PBS. The entire SPR sensing methodology was executed by a sequence of automatic procedure (baseline, association, dissociation, regeneration, and back to baseline steps). A sample solution containing a selected concentration of rH1N1HA in the PBS was injected in both channels from the 384 well microtiter plate and then association step was performed for 500 s and dissociation step was performed for

400 s followed by the regeneration of the sensor surface by addition of 10 mM HCl for 120 s. PBS (pH 7.5) was used as the running buffer solution throughout the study. For SPR sensing, an aliquot of the analyte solution (75 µL) was injected and mixed at 16.7 µL/s and this protocol was adopted with 1:12,800, 1:6,400, 1:3,200, 1:1,600, 1:800, and 1:400 dilution of rH1N1HA. The resultant data in these experiments were further utilized for the calculation kinetic parameters and also for plotting the calibration graph.

Optimization of Experimental Parameters

Effect of Temperature on SPR Response

Temperature is a well-known physical parameter to influence sensitivity of SPR measurement, and temperature variation allows to find out the temperature dependence of molecular interactions and also for optimizing detection conditions [39]. The determination of the reaction constants at different temperatures also gives the opportunity to access the thermodynamic parameters for anti-rH1N1HA and rH1N1HA interaction. The two-channel SPR system having a reference channel to compensate for bulk refractive index (RI) changes, non-specific binding, and minor temperature variations, while the temperature controller device (water bath) provides the temperature stability necessary for monitoring small changes in RI [40]. Changes in temperature affect the materials present on the SPR sensor chip and RI of aqueous sample solutions. Optimal temperature for binding of rH1N1HA (1:12,800 dilution) on immobilized anti-rH1N1HA was investigated by varying the temperature between 10 and 37 °C with a 3 °C increment.

Effect of pH on SPR Response

In order to know the effect of pH on the change in SPR angle, pH variation study was carried out using different buffers in the pH range from 4.0 to 9.0 with 0.5 increments. To maintain the pH, acetate buffer (pH 4.0 to 5.5), PBS buffer (pH 6.0 to 7.5), and glycine—NaOH buffer (pH 8.0 to 9.0) were used for the interaction of rH1N1HA with immobilized anti-rH1N1HA. The pH variation study was conducted from 4.0 to 9.0 as below pH 4.0 and above pH 9.0 protein losses its activity [38, 41].

Result and Discussion

Immobilization of Anti-rH1N1HA on 4-MBA Modified Gold SPR Sensor Chip

The most important requirement for the effective immobilization of ligand on the SPR gold chip surface is that the pH of

the ligand solution should be above pH 3.5 and below the 0.5 unit of isoelectric point of the ligand on the gold chip surface and so that the ligand carry opposite net charges. In addition, the electrostatic interactions involved in preconcentration are favored by low ionic strength in the coupling buffer [42]. Figure 1 represents the stepwise immobilization of anti-rH1N1HA on 4-MBA modified chip, and this process comprises of nine steps. Figure 1, in the first step, stabilization of baseline was performed for 120 s and the SPR angle observed for this process was -580 m° . In the second step, for chemical binding between the 4-MBA present on the gold SPR chip surface and free amino groups of anti-rH1N1HA, activation of carboxyl groups on 4-MBA modified gold SPR chip was carried out for 900 s with EDC-NHS which showed an angle change from -580 to -84 m° . In the third step, washing was conducted with PBS and the SPR angle shifted from -84 to -397.39 m° and this is due to the conversion of 4-MBA to its ester. In the fourth step, anti-rH1N1HA was injected on 4-MBA modified gold chip, allowed to interact 1800s and an increase in SPR angle is observed from -397.39 to 101.95 m° . In the fifth step, washing was performed, and this process resulted in the SPR angle change from 101.95 to 55.89 m° . The sixth step was performed to prevent non-specific binding and also for the blocking of unreacted NHS-ester groups on 4-MBA chip; 1,000 mM ethanolamine was used for blocking and allowed to react with sensor surface for 600 s; the angle change involved here was from 55.89 to 495.33 m° . In seventh step, washing was performed for 30 s as discussed in the experimental part which resulted in the SPR angle change from 495.33 to -104.46 m° . In the eighth step, regeneration was carried out with 0.1 M hydrochloric acid for 120 s and the

SPR angle change was from -104.46 to -94.65 m° . At last in ninth step, back to baseline process was conducted for 60 s, and the SPR angle change for this step was from -94.65 to 20.62 m° . The important process involved in Fig. 1 is also represented as Scheme 1.

SPR Characterization

The SPR angle of 4-MBA modified chip is shifted from -583 to 20.62 m° due to the covalent attachment of anti-rH1N1HA on 4-MBA adsorbed SPR chip results in a shift in the SPR angle, and this modification allows monitoring of the mass coverage (antigen interaction) at the surface with a high accuracy [43, 44]. Thus, the shift in SPR angle is attributed to the formation of thin layer of anti-rH1N1HA on 4-MBA modified gold SPR chip. The changes of SPR curve after anti-rH1N1HA immobilization on 4-MBA modified gold SPR chip is shown in Fig. 2. Angle change after ligand coupling observed is 454.52 m° on 4-MBA modified gold SPR chip after anti-rH1N1HA immobilization; this reveals the covalent attachment of 3.79 ng/mm^2 of anti-rH1N1HA on 4-MBA modified gold chip [45].

Electrochemical Impedance Spectroscopic Characterization

EIS was used to characterize the electron transfer properties of the 4-MBA modified SPR gold chip. In Fig. 3, the semicircle diameter at higher frequency corresponds to the charge transfer limited process (R_{ct}) [46]. In addition to the above, R_{ct} is most influential and direct parameter to reflect the in situ changes which are occurring on the modified chip/electrolyte interface. Hence, EIS studies were conducted, obtained the data, and fit, and simulation method was adopted to find out the R_{ct} value for bare gold SPR chip, modified gold SPR chip with 4-MBA after immobilization of anti-rH1N1HA and after interaction of rH1N1HA with anti-rH1N1HA immobilized 4-MBA modified gold SPR chip, the R_{ct} values are found to be 0.683, 390, and 535 $\text{k}\Omega$, respectively. Figure 3a is showing the EIS spectra for the bare gold SPR chip, Fig. 3b is showing EIS spectra for anti-rH1N1HA immobilized on 4-MBA modified gold SPR chip, and Fig. 3c is showing EIS spectra for after interaction of rH1N1HA with immobilized anti-rH1N1HA on 4-MBA modified gold SPR chip. As illustrated in Fig. 3, after immobilization of anti-rH1N1HA on 4-MBA, R_{ct} increases significantly, showing increase in the semicircle diameter on the Nyquist plot. This is due to the presence of anti-rH1N1HA and 4-MBA on gold SPR chip surface [47]. It is observed from Fig. 3c that the Nyquist diagram includes a squeezed semicircle portion at higher frequencies, which corresponds to the electron transfer limited process, followed by a linear part characteristic of the lower frequency attributable to a diffusion-limited process [48].

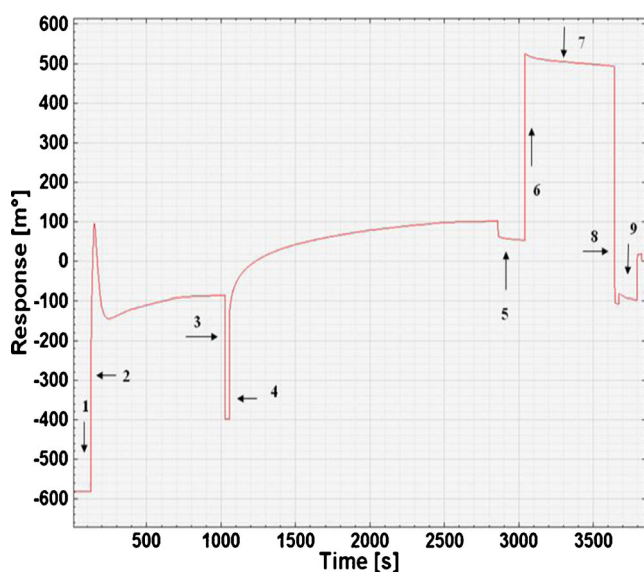
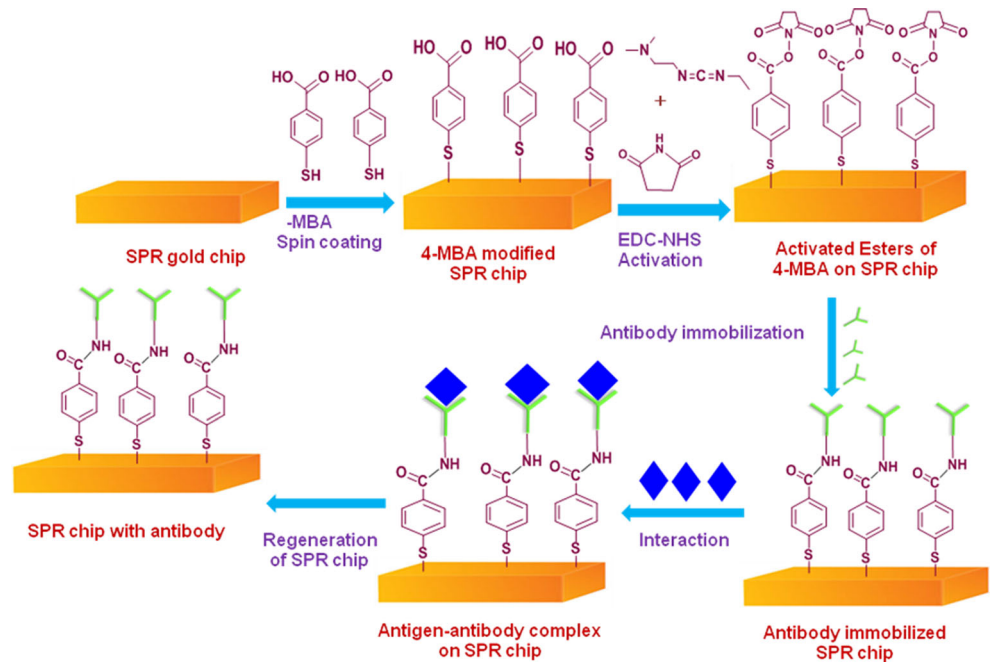


Fig. 1 Sensorgram showing different steps (1 baseline, 2 EDC-NHS activation, 3 washing, 4 anti-rH1N1HA coupling, 5 washing, 6 deactivation, 7 washing, 8 regeneration, and 9 back to baseline) involved in the immobilization of anti-rH1N1HA on 4-MBA modified gold chip

Scheme 1 Different steps involved in the immobilization of anti-rH1N1HA on 4-MBA modified gold SPR chip and its application for sensing



Interaction of rH1N1HA With the Immobilized Anti-rH1N1HA on Gold SPR Chip

The anti-rH1N1HA immobilized on the SPR sensor chip was utilized for the sensing of different concentrations of rH1N1HA as discussed in the experimental part, and the results are depicted as SPR sensorgram in Fig. 4, and this figure exhibit concentration dependant angle changes with

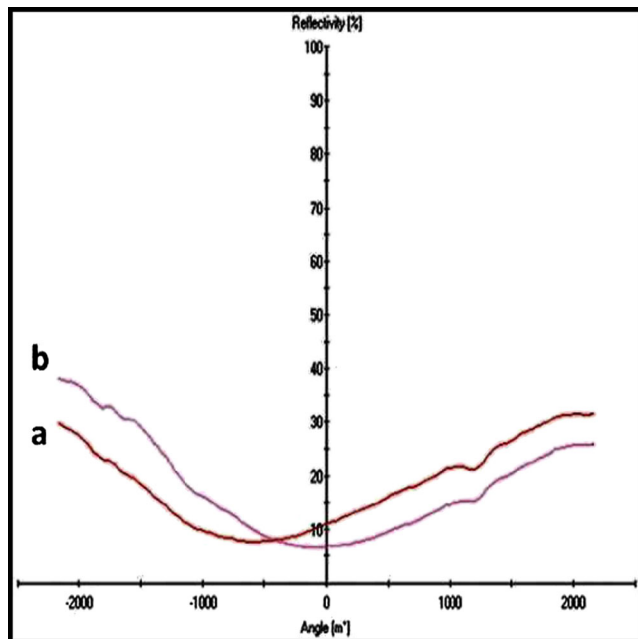


Fig. 2 SPR angle shift curve for *a* 4-MBA modified gold SPR chip and *b* after immobilization of anti-rH1N1HA with 4-MBA modified gold SPR chip

various concentrations of rH1N1HA from 0.015 pg to 1.5 ng/ml. Calibration graph was constructed using the response of angle changes (Fig. 4) against log concentration of antigen and presented in Fig. 5. The limit of detection (LOD) of the present method was calculated experimentally and was found to be 0.015 pg/ml rH1N1HA, and this is the minimum concentration which showed the response during the interaction with the immobilized anti-rH1N1HA on 4-MBA modified chip. It is noteworthy to compare here the result of the present work with our previous report [34], where we have

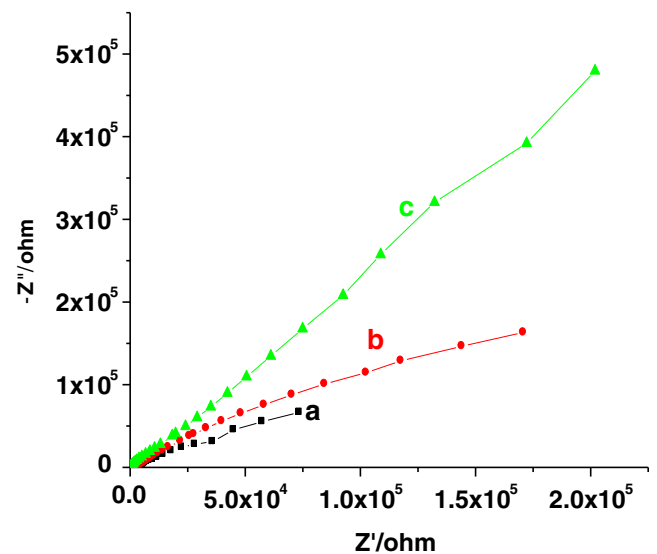


Fig. 3 EIS for *a* bare gold chip, *b* 4-MBA modified gold chip after immobilization of anti-rH1N1HA, and *c* after interaction of rH1N1HA with anti-rH1N1HA 4-MBA modified gold chip

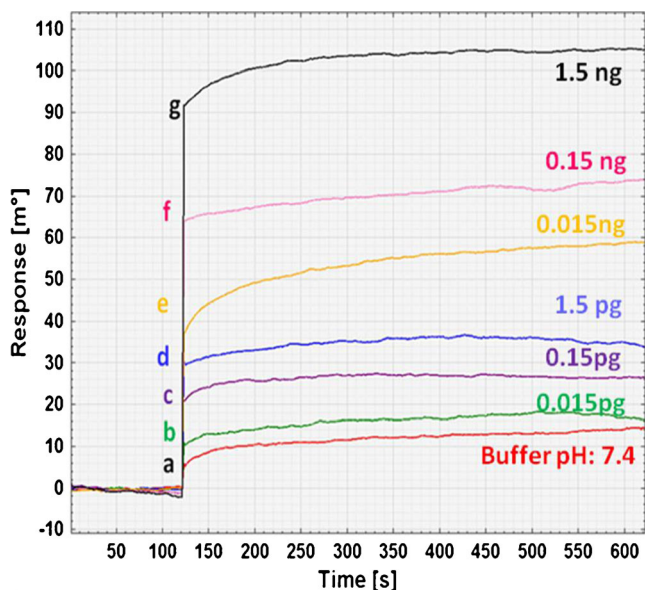


Fig. 4 SPR sensor response for the interaction of different dilution of rH1N1HA (a buffer pH 7.5, b 1:12,800, c 1:6,400, d 1:3,200, e 1:1,600, f 1:800, and g 1:400) with immobilized anti-rH1N1HA

immobilized the rH1N1HA on 4-MBA modified SPR chip and detected anti-rH1N1HA; the LOD found for that interaction was 0.5 ng/ml, and this value is very less than that of the value obtained in the present study 0.015 pg/ml. The findings of the present study and our earlier report [34] implies that SPR detection of rH1N1HA using anti-rH1N1HA immobilized SPR chip is better and more sensitive than that of the SPR detection of anti-rH1N1HA using rH1N1HA immobilized SPR chip (reason for this observation is discussed in Evaluation of kinetics involved in between the interaction of rH1N1HA with immobilized anti-rH1N1HA below).

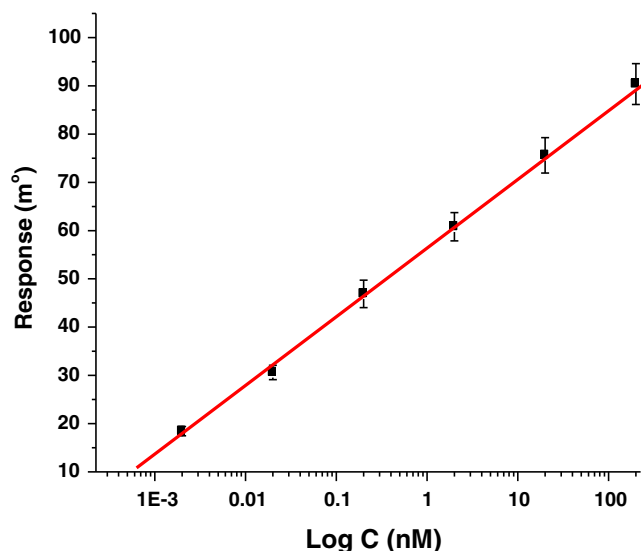


Fig. 5 Calibration graph for the response of angle changes against log concentration of rH1N1HA with immobilized anti-rH1N1HA; temperature 25 °C and pH 7.5

Evaluation of Kinetics Involved in Between the Interaction of rH1N1HA With Immobilized Anti-rH1N1HA

The affinity interactions between immobilized antibody and antigen were characterized by the equilibrium constant (K_D). The data were fitted using a simple 1:1 interaction model [49]. $A + B = AB$, where “A” is the injected analyte, “B” is the immobilized ligand, and “AB” is the analyte–ligand complex formed during the interaction process. In the SPR system, the signal R is proportional to the amount of (AB) and the R_{max} is proportional to the initial (B). Hence, in this study, kinetic parameters such as K_D and B_{max} value were calculated for binding of rH1N1HA with immobilized anti-rH1N1HA using the software and were found to be 0.278 pM and 84.51 m° , respectively. This low K_D value (if $K_D \leq 10$ nM, then it represents the high affinity interactions) represents high affinity interaction of the antigen with the immobilized anti-rH1N1HA [50]. This K_D value was compared with the K_D value of our earlier report [34], where we have immobilized the rH1N1HA on 4-MBA modified SPR chip and interacted anti-rH1N1HA; the K_D value found for that interaction was 49.7 nM, and that value is very less than that of the value obtained in the present study 0.278 pM. These findings (present study and our earlier report [34]) implies that SPR interaction of rH1N1HA with anti-rH1N1HA immobilized SPR chip is having high affinity and the SPR interaction of anti-rH1N1HA with rH1N1HA immobilized SPR chip is having low affinity based on the K_D values. This is the reason for the good LOD obtained in the present study for the SPR detection of rH1N1HA.

Evaluation of Thermodynamic Parameters

The K_D obtained from kinetic evaluation software for the binding of rH1N1HA with immobilized anti-rH1N1HA was further exploited first time to calculate the thermodynamic parameters such as change in Gibb’s free energy (ΔG), change in enthalpy (ΔH), and change in entropy (ΔS) associated with binding using Van’t Hoff equation [51, 52]:

$$\Delta G = -RT \ln K_A = \Delta H - T\Delta S \tag{1}$$

$$\Delta H = R \frac{T_2 T_1}{T_2 - T_1} \frac{\ln K_2}{K_1} \tag{2}$$

$$K_A = 1/K_D \tag{3}$$

where R indicates universal gas constant, K_A affinity constant, T temperature, and K_1 and K_2 are affinity constants for association at T_1 and T_2 temperature, respectively. ΔG , ΔH ,

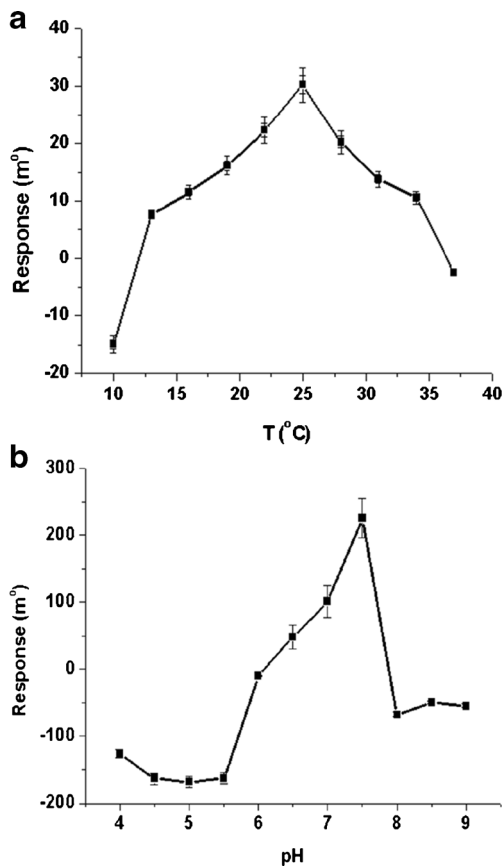


Fig. 6 **a** Effect of temperature on the interaction of rH1N1HA with the immobilized anti-rH1N1HA on 4-MBA modified gold chip. Temperature range 10 °C to 37 °C, pH 7.5 and rH1N1HA (dilution 1:12,800). **b** Effect of pH on the interaction of rH1N1HA with the immobilized anti-rH1N1HA; pH range 4.0 to 9.0, temperature 25 °C and rH1N1HA (dilution 1:12,800)

and ΔS are change in Gibb's free energy, change in the enthalpy and change in the entropy due to the interaction of rH1N1HA with immobilized anti-rH1N1HA.

The value of the change in Gibb's free energy for binding of rH1N1HA with immobilized anti-rH1N1HA was found to be -49.78 kJ/mol at 298 K. The negative value of ΔG indicates the spontaneous interaction of rH1N1HA with immobilized anti-rH1N1HA. The calculated value of ΔH using Van't Hoff wizard in kinetic evaluation software was found to be 98.25 kcal/mol. The positive value of ΔH reveals the interaction of rH1N1HA with immobilized anti-rH1N1HA as endothermic process [53]. The value of ΔS for binding of rH1N1HA with immobilized anti-rH1N1HA was calculated using software and was found to be 72.59 cal/mol.K. The observed positive value in entropy indicates that the interaction can be explained by Langmuir replacement reaction that exhibits Langmuir type isotherm [51]. Langmuir replacement reaction suggests that the observed positive entropy is because of desorption of water molecules from either antibody or antigen or both. The significance of desorption of water

molecules from protein surfaces during ligand-protein binding was reported earlier [54].

Effect of Temperature and pH

Temperature variation study was carried out in order to know the effect of temperature on SPR response during the interaction of rH1N1HA with immobilized anti-rH1N1HA. Upon increasing temperature from 10 to 25 °C, an increase in SPR angle is observed as shown in Fig. 6a, and beyond 25 °C, SPR angle decreased. Hence, 25 °C is used as optimum temperature for the interaction of rH1N1HA with its immobilized anti-rH1N1HA on 4-MBA chip [55, 56]. Figure 6b shows the effect of pH on SPR angle due to the interaction of rH1N1HA with its immobilized anti-rH1N1HA. It is observed from Fig. 6b that SPR angle increased as increase in pH up to 7.5 and then decreased up to pH 9.0. This observation is probably due to the pH-dependent structural changes and electrostatic interactions on the sensor disc between antibody and antigen as reported earlier [57]. All these observations suggests that in pH 7.5 PBS buffer, rH1N1HA, and anti-rH1N1HA interaction is more effective and thereby resulted in more angle change; hence, pH 7.5 was preferred in this study.

Conclusion

A label-free, real-time SPR detection methodology for rH1N1HA was developed using a 4-MBA modified SPR gold chip. The modification of gold chip with 4-MBA and anti-rH1N1HA was confirmed in situ by SPR and EIS. The detection time for rH1N1HA by SPR is found to be less than 10 min. By using kinetic evaluation software, K_D and B_{max} values were calculated and found to be 0.278 pM and 84.51 m°, respectively. Moreover, thermodynamic parameters such as ΔG , ΔH , and ΔS were determined first time for the interaction between rH1N1HA and anti-rH1N1HA, and the values revealed that this interaction is spontaneous, endothermic, and entropy-driven one. This study also indicates that rH1N1HA interaction with anti-rH1N1HA immobilized SPR chip is having high affinity when compared to anti-rH1N1HA interaction with rH1N1HA immobilized SPR chip based on K_D and thermodynamic parameters. The present SPR-based approach may used in the hospitals equipped with SPR for the early screening of clinical samples in a label-free real-time manner.

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