Use of a Quartz Crystal Microbalance for Studying the Effects of Propyl Gallate on the Interactions of Interleukin-6 and Its Soluble Receptor

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Abstract

Quartz Crystal Microbalance (QCM) is known as a surface sensitive analytical device capable of on-line monitoring of interfacial reactions. To study the interaction between interleukin-6 (IL-6) and its soluble receptor (sIL-6R), as well as the effects of propyl gallate on IL-6/sIL-6R binding based on it, a QCM was employed to monitor the binding responses. In the experiments, sIL-6R was immobilized on the surface of QCM substrate, then different concentrations of IL-6 were added onto the substrate, respectively. The response frequency shift caused by the binding between IL-6 and sIL-6R was recorded. The results showed that sIL-6R was immobilized on the substrate surface with the density of 3.43 ng/mm². IL-6 then bound to the immobilized sIL-6R specifically. FAST fit program was used to analyse the data obtained of the association of IL-6 and immobilized sIL-6R, fitting it to the biphasic kinetic equations. Kinetic constants for IL-6 binding to sIL-6R were determined from the plots of Kd versus the concentration of IL-6, kd = 3.41×10⁴ M⁻¹ s⁻¹, kₐ = 1.80×10⁻² s⁻¹, thus Kd = 3.41×10⁻¹ M⁻¹. Furthermore, the effects of propyl gallate (PG) on interaction of IL-6 with sIL-6R were studied. The results showed that PG was able to enhance the binding between IL-6 and sIL-6R. We propose that the interaction between IL-6 and sIL-6R is one of the targets which PG can act on in vivo. The QCM offers an effective way and a new clue to the study on the mechanism of propyl gallate at molecular level.

Keywords: Interleukin-6 (IL-6); Soluble interleukin-6 receptor (sIL-6R); Propyl gallate; QCM; Interaction; Kinetic analysis

Introduction

Quartz crystal microbalance (QCM) is known independently as a surface sensitive analytical device capable of on-line monitoring of interfacial reactions and thin film depositions [1,2]. QCM measures changes in acoustic thickness or mechanical resonance properties of a thin film deposited on a metal electrode (e.g. gold, silver, and copper, etc.) [3,4]. As a surface analytical tool, combined QCM data collection and analysis allow one to take the advantage of the strengths, to test the validity of the assumptions and to gain a better understanding of a specific interfacial reaction. QCM has been widely used for biological analysis, clinical diagnosis, and environmental monitoring [5,6]. The liquid cell configuration of this device makes it suitable for real-time study of bioaffinity reactions in relevant solution conditions of temperature, flow rate, pH and ionic strength. In the recent years, the applications of QCM for biological analyses have been reported increasingly, including immunoassay of BSA [7], enterotoxin detection [8], enzymatic analysis for hydrogen peroxide quantification [9], and blood plasma coagulation determination [10].

PG is a well-known synthetic antioxidant on the US Food and Drug Administration list (Figure 1). It is a derivative of gallic acid which is one of the effective ingredients of commonly used traditional Chinese medicine (TCM) Paeoniae Radix. Besides its antioxidant activity, PG has many pharmacological effects such as free radical elimination [11], inhibiting platelet aggregation [12,13], anti-inflammatory effects [14], and anti-tumor activities [15]. It is widely used to nourish blood, activate circulation, alleviate pain, regulate menstruation, treat liver disease and cancer. But until now, the researches on PG have been focused mainly on its curative effects at the holistic level. There are no detailed studies on the mechanisms and dynamics of action at the molecular level, which limits the further clinical use of PG.

Interleukin-6 (IL-6) is a multifunctional protein that mediates both immune and inflammatory responses. Recombinant human IL-6 is a kind of polypeptide containing 184 amino acid residues whose molecular weight is estimated to be 20.9 kDa [16]. IL-6 is secreted by many cell types, activates monocytes, macrophages, T cells, endothelial cells and bone marrow stroma cells [17]. IL-6 is a cytokine with many functions, which induces the transformation of activated B cells into immunoglobulin-secreting cells and the production of acute phase proteins by hepatocytes [18]. IL-6 acts on target cells via a receptor consisting of two transmembrane glycoproteins, gp80 and gp130. The binding of IL-6 to its specific receptor (gp80, IL-6R) triggers the dimerization of the signal-transducing receptor subunit gp130 [19].

Figure 1: Chemical structure of propyl gallate, Mr 213.
The soluble form of the IL-6R (sIL-6R) when linked to IL-6 triggers the dimerization of gp130 and in contrast to several other soluble cytokine receptors, which inhibit ligand function. In fact, the IL-6/sIL-6R complex rather than IL-6 is believed to be the active form in vivo [20]. The soluble IL-6/sIL-6R complex can activate signal transduction and biological responses in cells that possess gp130 and do not express gp80 IL-6R [21,22]. IL-6R produced at sites of inflammation or in the peripheral blood in response to stimuli released by inflammatory sites (IL-1 or TNFα) binds the serum sIL-6R and is transferred in the blood as an IL-6/sIL-6R complex [23]. This binding protects IL-6 from protease degradation and stabilizes its bioactivity. The routine method to study on the interaction between IL-6 and sIL-6R is enzyme-linked immunodeficient assay (ELISA), which is complicated, time-cost, only detects the equilibrium binding values and cannot monitor the process of the binding in real time.

In the present paper, a QCM was employed to monitor the interaction between IL-6 and sIL-6R. Moreover, to evaluate the affinity of the binding, the interactions between sIL-6R and IL-6 with series of concentrations were detected and the kinetic analyses between IL-6 and sIL-6R were carried out using FAST fit program. Furthermore, the effects of PG on IL-6/sIL-6R binding based on the interaction analysis are observed to find out the molecular mechanism of propyl gallate.

Experimental

Materials

N-hydroxysuccinimide(NHS), 1-ethyl-3-(3-dimethylaminopropyl) carbodiimide (EDC), ethanolamine and PG were purchased from Sigma-Aldrich (St Louis, MO, USA). Recombinant human Interleukin-6 (IL-6) and recombinant human Interleukin-6 soluble receptor (sIL-6R) were purchased from PeproTech EC (London, UK). AT-cut 9 MHz QCM quartz crystal substrates were purchased from SEIKO EG&G. Phosphate-buffered saline Tween-20 (PBST, pH7.4) was composed of 0.01 M Na₂HPO₄/NaH₂PO₄, 0.138 M NaCl, 0.0027 M KCl and 0.05% Tween-20. All solutions were made by using deionized water. This process was repeated three more times.

Preparation of AT-cut 9 MHz QCM quartz crystal substrate

The QCM system employed in this study consists of an AT-cut 9 MHz quartz crystal substrate. Five V dc was applied to the circuit to drive the crystal substrate and the frequency was monitored with a frequency counter (Model QCA917, SEIKO EG&G) connected to a computer. The quartz crystal was employed as a working electrode. The application of a small change in mass ∆m of the gold substrate is detected and the kinetic analyses between IL-6 and sIL-6R were carried out using FAST fit program. Furthermore, the gold substrate was cleaned by using acetone for 30 min and incubated in anhydrous methanol for 30 min. By the addition of 20 μl of α-TA solution on the surface of the substrate for 50 min. Disulfides are known as a kind of molecules to form a stable self-assembled monolayer (SAM) on a gold surface owing to the strong S–Au covalent bond [29]. Then a SAM of α-TA was formed on the gold substrate. After the treated substrate was rinsed with deionized water for 3 times, α-TA treatment surface introducing COOH groups made possible to immobilize avidin through the formation of amide bonds. EDC/NHS mixture was dropped to the SAM of α-TA on the surface of the gold substrate. The function of the EDC/NHS is to activate and promote the formation of the covalent linkages by forming the N-hydroxysuccinimide ester (Scheme 1).

Then different concentrations of sIL-6R (10 μl) were added onto the α-TA-bound surface of the gold substrate for 50 min at room temperature. The carboxyl on the SAM of α-TA bound NHS under the action of EDC and created active ester intermediate which bound the primary amines of the protein to make amide bond. Thus, sIL-6R was covalently immobilized on the gold substrate. The optimum concentration of sIL-6R is discussed later. Then the substrate was rinsed thoroughly with deionized water (Scheme 1).

Detection of the binding between sIL-6R and Paeoniae IL-6

Unreacted NHS-esters were blocked by washing with 1 M ethanolamine, pH 8.5. Ten 10 μl of IL-6 solution was added onto the surface of the gold substrate modified by sIL-6R, equilibrating for 20 min. The frequency response was recorded. The amount of the immobilized IL-6 was determined by the equation (1). The progressed binding of IL-6 on the sIL-6R-bound surface of the gold substrate is schematically depicted in Scheme 1.
Kinetic Analysis of the Binding between IL-6 and sIL-6R: To analyze the kinetics of binding of IL-6 to sIL-6R, determination of kinetic and equilibrium constants was achieved by adding a wide range of ligate concentrations (in the ng-μg ml⁻¹ range) and allowing association to occur. Equilibrate sensor substrate for several minutes in 40 μl of PBST, added 10 μl of IL-6 solution into the cuvette, 10 min so association was recorded. Remove the IL-6 solution by washing the substrate 3 times with 50 μl of PBST. Regenerate the bound IL-6 by washing the substrate in 50 μl of 10 mM HCl, until the response signal returned to baseline. Wash sufficiently with PBST for re-use. Each binding cycle with different concentrations of IL-6 was analyzed as described above. Data obtained was analyzed using FAST fit program and K_eq was calculated.

Effects of Propyl Gallate on the Binding between IL-6 and sIL-6R: After three washes with PBST, 10 μl of IL-6 (0.01 mg ml⁻¹), preincubated for 15min with propyl gallate concentration of 500 nM, was added into the sIL-6R modified substrate. Then, detecte the binding between IL-6 and sIL-6R as described above.

Results and Discussion

sIL-6R Immobilization on the Gold Substrates of QCM

Figure 2 shows the response of a QCM immobilized by sIL-6R and IL-6 binding on the surface of the gold substrate. Response (1) is at the equilibration and a stabilized baseline after the addition of PBS. Response (2) corresponds to the addition of sIL-6R. Response (2) to (3) is buffer washing to remove unreacted sIL-6R. Response (4) is the blocking of non-coupled activated NHS-esters with ethanolamine. Response (5) is buffer washing to remove ethanolamine. Response (6) is the addition of IL-6. Response (7) is buffer washing to remove unreacted IL-6 (8) corresponds to ∆F.

Response (1) is at the equilibration and a stabilized baseline after the addition of PBS. Response (2) to (3) is buffer washing to remove unreacted sIL-6R. Response (4) is the blocking of non-coupled activated NHS-esters with ethanolamine. Response (5) is buffer washing to remove ethanolamine. Response (7) is buffer washing to remove unreacted IL-6 (8) corresponds to ∆F.

The mass of the gold substrate increased with the addition of α-TA on the surface of the gold substrate. The resonance frequency of QCM was simultaneously dropped. The gold substrate was rinsed with deionized water in order to remove the adsorbed species. The sIL-6R addition led a shift of response frequency to decreasing direction (2)-(4), indicating a time-dependent accumulation of the IL-6 molecules on the surface. The adsorbed values of IL-6 on the substrate were calculated as 130.919 ng cm⁻². The results mentioned above are concluded that IL-6 binds sIL-6R.

The specific binding between IL-6 and sIL-6R was evaluated by a blank control experiment without anchoring sIL-6R on the substrate surface. No detectable binding signal was obtained, showing that IL-6 binds sIL-6R specifically on the substrate surface.

Kinetic analysis of the binding between IL-6 and sIL-6R

Figure 3 shows the association curve of IL-6 with different concentrations binding to immobilized sIL-6R. As IL-6 concentration increased, the initial binding velocity increased and the time taken to approach equilibrium decreased.

Data is analyzed using the FAST fit program, specifically designed for biomolecular interaction analysis using the IAsys optical biosensor. It performs the detailed mathematical analysis of the association and dissociation curves obtained in QCM. By specifying IL-6 concentration and the region to analyze and fitting the experimental data for each IL-6 concentration to corresponding equations, the association curves showed good fitting to double phase association.

Each correctly-fitted association curve provides a value for k_on, the apparent association rate constant, which was used to determine k_on (association rate constant) and k_off (dissociation rate constant), as described by:

\[ k_{on} = k_{ass} [L] + k_{off} \]
these $k_{\text{on}}$ values are obtained for each concentration of ligate, IL-6. The certain option in FAST fit is used to produce a plot of $k_{\text{on}}$ versus IL-6 concentration (Figure 4).

The gradient of the line in Figure 4 is used to calculate $k_{\text{on}}$, $k_{\text{off}}$, which is obtained from the intercept of Figure 3. Thus $k_{\text{on}}=3.41 \times 10^4 \text{ M}^{-1} \text{s}^{-1}$, $k_{\text{off}}=1.80 \times 10^3 \text{ s}^{-1}$, $K_D$ may be determined from the relationship:

$$K_D = \frac{k_{\text{off}}}{k_{\text{on}}}$$

which indicated that sIL-6R bind IL-6 with high affinity. Compared to the $K_D$ of IL-6/sIL-6R binding, reported as $\sim 10^{-7} \text{ M}$ [27], it can be inferred that the binding ability of sIL-6R to IL-6 is lower than sIL-6R. The results indicate that sIL-6R binding ability to IL-6 is decreased, which is probably attributable to the denaturation that occurred when sIL-6R was produced. Furthermore, the developed approach showed very desirable reproducibility. The relative standard deviations (RSDs) of $K_D$ are 2.84%, 3.17%, 2.72%, respectively, in three repetitive assays of 80 nM, 320 nM, 1280 nM of TNF-α. These data manifested that the developed approach held great potential with excellent reproducibility.

**Effects of Propyl Gallate on the Binding between IL-6 and sIL-6R**

Figure 5 showed the experimental results of the binding reactions of IL-6 with PG, as well as without PG, on the immobilized sIL-6R surface. As PG involved, the initial velocity of IL-6/sIL-6R binding increased and the time taken to approach equilibrium decreased, which indicated that PG can also enhance the binding of IL-6 to sIL-6R.

**Conclusions**

In the present study, based on the interaction analyses between IL-6 and sIL-6R, the effects of PG on IL-6/sIL-6R binding were studied by using QCM. The results showed that IL-6 can bind sIL-6R specifically in a dose-dependent manner. Data achieved was analyzed using the FAST fit program, $K_D$ value is $5.28 \times 10^{-7} \text{ M}$, indicated that sIL-6R binds IL-6 with high affinity. Thus sIL-6R may protect cell and tissue form the pathologic injury of IL-6 by competing the cell-surface receptor and neutralizing IL-6 bioactivity, which agree with previous reports by others that the administration of sIL-6R can prevent the adverse pathologic sequelae of exaggerated IL-6 production [30].

Moreover, the acquisition of kinetic analyses will be of benefit for obtaining a better insight on the study on IL-6/sIL-6R system and further studies based on it. In this work, the effects of propyl gallate on interaction between IL-6 and sIL-6R were observed. The experimental results showed that PG enhanced IL-6/sIL-6R binding. Though the mechanism of PG is not clear and the roles of IL-6 and sIL-6R are debated, it can be concluded that the binding between IL-6 and sIL-6R is another target which PG can act on in vivo, that is, PG can enhance the cellular protection of sIL-6R through the increase neutralization to IL-6, which is in agreement with the experimental and clinical results reported [31].

A QCM was employed to explore the mechanisms of PG for the first time. This finding offers an effective way to study the biomolecular interaction as well as a new clue to the study of the mechanisms of PG based on cytokine/receptor binding. Though various methods are useful for studying the interaction process between cytokines and receptors, many of these methods require labeling of one of the two biomolecules with a fluorescent or radioactive tag and a long time to equilibrium. QCM appeared to be a high sensitive instrument for monitoring and quantifying the binding of the cytokine to the corresponding receptors, which is fundamental for obtaining a better insight on the molecular interactions between the cytokine and receptor involved in various of physiological and pathophysiological processes. Furthermore there are other advantages of this new method, such as that the amount of both ligand and ligate needed to obtain informative results is very low, the time required to perform an assay is very short and all the procedures are label free. It has tremendous application prospect in the studies on the molecular mechanisms of TCM related to the cytokine/receptor interaction. Further studies in this area are in progress.

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**References**


