

Original Research Paper

Binding Mechanism of Bovine Serum Albumin to Troxerutin by Synchronous Fluorescence Spectroscopy and Chemometrics

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Abstract: Previous works have reported the binding of Bovine Serum Albumin (BSA) to troxerutin by absorption and fluorescence spectroscopy. In order to extend the investigation of their binding characteristics, the study presented here was designed to evaluate their interaction by the synchronous fluorescence spectroscopy and Multivariate Curve Resolution With Alternating Least Squares (MCR-ALS) under simulative physiological conditions. The Evolving Factor Analysis (EFA) and MCR-ALS analysis results show that there are three chemical species in the troxerutin-BSA system, which reveal that the fluorescence quenching of BSA caused by troxerutin may be a static quenching mechanism. The spectral profile of each compound was obtained by the soft of MCR-ALS. The obtained average binding constant is $6.43 \times 10^5 \text{ mol}^{-1} \text{ L}$ for BSA and troxerutin concentration ratio is 7 while that is $6.89 \times 10^5 \text{ mol}^{-1} \text{ L}$ for troxerutin and BSA concentration ratio is 5. The study might further help us to better understand the structural features and toxicological action of troxerutin at the molecular level.

Keywords: Bovine Serum Albumin (BSA), Troxerutin, Synchronous Fluorescence Spectroscopy, MCR-ALS

Introduction

Troxerutin is a flavonoid derivative, could be extracted from coffee, tea, cereal grains and a variety of fruits (Marzin *et al.*, 1987). It exerts a great impact on inhibiting the role of red blood cells and platelet aggregation, preventing thrombosis, increasing the content of oxygen in the blood and improving micro-circulation etc (Adam *et al.*, 2005). Thank for its pharmaceutical properties, it has been widely used to treat the arteriosclerosis, thrombophlebitis, edema caused by increased vascular permeability, Chronic Venous Insufficiency (CVI) (Pentz *et al.*, 2005). So far, considerable works have been focused on the pharmaceutical properties of troxerutin in order to further to understand its biological activity (Lu *et al.*, 2011; Fan *et al.*, 2009; Maurya *et al.*, 2005). Subastri *et al.* (2015) reported its interaction with transfer RNA and revealed that it may have potential therapeutic application. Heidarzadeh *et al.* (2014) showed that troxerutin may reduce tissue injury and lipid

peroxidation. Wang *et al.* (2009) studied the binding of troxerutin to Bovine Serum Albumin (BSA) and revealed that the fluorescence quenching of BSA caused by troxerutin is a static quenching procedure.

Serum albumins are the most abundant protein constituents of blood plasma, which are responsible for maintaining the pH in blood and transporting drugs in living organisms (Anbazhagan and Renganathan, 2008; Kragh-Hansen, 1981). Among the serum albumins, BSA has been widely used in biophysical studies to clarifying the binding mechanism of small molecules to proteins due to its physiological functions and 76% of structural similarity with Human Serum Albumin (HSA) (Sulkowska, 2002; Dasmandal *et al.*, 2015). Investigation on binding of drugs to serum albumins may help us to understand drug's distribution and toxicity in the biological systems.

In recent years, several optical techniques have been carried out to clarify the interaction mechanism of drugs with serum albumins, such as UV-vis and fluorescence

spectroscopy, Raman spectroscopy and Circular Dichroism (CD) and so on. Among these methods, the spectrophotometry is the most common and convincing methods (Cui *et al.*, 2004; Lin and Koenig, 1976; Gelamo *et al.*, 2002). With the development of chemometrics algorithms and high precision instruments, multi-dimensional data of complex reaction system could be easily obtained. MCR-ALS is a very powerful tool to handle deviations from trilinearity because it is easy to deal with the data and also, it is a powerful chemometrics tool (Ni *et al.*, 2011; Azzouz and Tauler, 2008).

In the previous study, we have reported the binding mechanism of troxerutin to BSA by spectral method involving fluorescence spectroscopy (Wang *et al.*, 2009). The binding distance, binding site, binding constants and the effect of troxerutin on the conformation of BSA were studied. In order to extend to the investigation of their interactions, the study presented here was designed by using the synchronous fluorescence spectroscopy and MCR-ALS under simulative physiological conditions. The number of chemical species that present in the interaction system of troxerutin with BSA was analyzed by using Evolving Factor Analysis (EFA) based on synchronous fluorescence spectroscopy data. Subsequently, the spectral profile of each compound was achieved by a soft of MCR-ALS.

Materials and Methods

Materials and Apparatus

Troxerutin was procured from Shanxi Yabao of China. The troxerutin solution (3.0×10^{-3} mol L⁻¹) was prepared in phosphate buffer solution of pH 7.4. BSA was obtained from Boquan Biochemical Technology of China and has been used without further purification. The stock solution of BSA (3.0×10^{-5} mol L⁻¹) was prepared in phosphate buffer solution of pH 7.4 which containing 0.1 mol L⁻¹ NaCl. The synchronous spectra were recorded by spectrofluorophotometer of RF-5301PC (Shimadzu, Japan) equipped with a quartz cell of 1.0 cm at room temperature. The slit widths of emission and excitation were used in 5.0 nm.

Experimental Procedure

Experiment 1: The BSA concentration was kept at 1.0×10^{-6} mol L⁻¹, then, different concentration range (0.0 - 7.5×10^{-6} mol L⁻¹ at intervals 0.5×10^{-6} mol L⁻¹, total 16 solutions) of troxerutin was added.

Experiment 2: The troxerutin concentration was kept at 1.0×10^{-6} mol L⁻¹, then, different concentration range (0.0 - 7.5×10^{-6} mol L⁻¹ at intervals 0.5×10^{-6} mol L⁻¹, total 16 solutions) of BSA was added.

Evolving Factor Analysis (EFA)

The Evolving Factor Analysis (EFA) is a chemometric technique of identifying the nature structure of multivariate observation and treating dimension reduction, has been used to analyze the number of chemical species (Amrhein *et al.*, 1996; Maeder, 1987). EFA is calculated from two directions, namely forward direction and reverse direction. The eigenvalue of the X_i sub array can be calculated by the following formula:

$$X_i = YU^T \quad (1)$$

where, Y ($I \times N$) is the score matrix, U^T ($N \times N$) is the loading matrix, N is the number of pure substance, X_i is composed of the $i = 1, 2, \dots$ line of the data matrix X .

The main idea of EFA is to follow the evolution or change of the rank of Y with progressing elution by rank analysis of the sub matrices Y . The appearance of each new component at the detector is intrinsically associated with the increase of the rank by one. Obviously there has to be a minimal difference in the retention times and, further, the spectra data have to be linearly independent.

Multivariate Curve Resolution-Alternating Least Squares (MCR-ALS)

The soft of MCR-ALS, a recently developed method, could evaluate the concentration and spectral profile of the chemical species for multivariate data (Maeder and Zilian, 1988).

In order to obtain the concentration profile and pure spectra of spectroscopic ally active components in the drug-protein system, the soft of MCR-ALS procedure was adopted to analyze the experimental data matrix D ($N_C \times N_S$). The D can be decomposed according to the following equation (Jaumot *et al.*, 2005):

$$D = CS^T + E \quad (2)$$

where, C and S^T is the concentration profiles with the dimension of $N_C \times N$ and spectral profiles with the dimension of $N \times N_S$; respectively. E ($N_C \times N_S$) is the residual matrix containing the data variance unexplained by CS^T . N_C is the number of spectra recorded at different concentration ratio of ([BSA]: [Troxerutin]), N is the number of the detected chemical species and N_S is the number of the wavelengths.

In order to eliminate the uncertainties of rotation and solve the problem of low matrix resolution, an extended matrix was used in this study. For this purpose, two kinds of experiments were designed to obtain the data matrix D^{BSA} and $D^{Troxerutin}$ (see experiment 1). The extended matrix can be obtained from the synchronous spectroscopy data matrix at two modes of experiment 1 and 2, which are overlapped in the column direction. The

rows of new matrix are the sum of two rows of a single matrix. The extended matrix can be expressed as Equation 3:

$$\begin{bmatrix} D^{BSA} \\ D^{Troxeutin} \end{bmatrix} = \begin{bmatrix} C^{BSA} \\ C^{Troxeutin} \end{bmatrix} S^T + \begin{bmatrix} E^{BSA} \\ E^{Troxeutin} \end{bmatrix} \quad (3)$$

where, $[D^{BSA}, D^{Troxeutin}]$ is the column-wise augmented matrix. The $D^{Troxeutin}$ and D^{BSA} , correspond to two different experiments, 1 and 2, respectively (see experiment 2).

Results and Discussion

The Synchronous Fluorescence Spectroscopy of Troxeutin-BSA Interaction

Figure 1 and 2 show the synchronous fluorescence spectroscopy of the binding of troxeutin to BSA in the two different experiments 1 and 2, in which $\Delta\lambda$ is equal to 20 nm, under simulative physiological conditions. According to Fig. 1, the synchronous fluorescence intensity of BSA gradually decreased with the increasing the troxeutin concentration, which may indicate that the binding of troxeutin to BSA had occurred as reported in earlier work. Further, the maximum emission wavelength of BSA show a slight blue shift, which indicates that the micro-environment around of BSA has been changed (Singh *et al.*, 2015; Tao *et al.*, 2015). From Fig. 2, one can see that the synchronous fluorescence intensity of troxeutin gradually increased with increasing the BSA concentration.

However, it is very difficult to confirm whether the reaction has occurred between troxeutin and BSA and cannot directly determine that the quenching of spectroscopy is fluorescence quenching or spectral superposition. In order to further study the binding of troxeutin to BSA, the synchronous fluorescence spectroscopy data was studied by combination of EFA and MCR-ALS methods.

The Chemical Species Analysis

Figure 3 shows the EFA plots of the synchronous spectroscopy between troxeutin and BSA. The blue horizontal line was retrieved by related noise and EFA. According to (Amrhein *et al.*, 1996), the number of significant factors containing the spectral evolution recorded degradation process, which is equal to the number of chemical species contributing to the analyzed spectra or the number of independent reactions +1. In the forward analysis, the first degradation variability of spectral data can be explained by two factors. In the beginning of the reaction, the presence of two factors must be corresponding to two chemical species of the process, in which BSA is degraded. The ANOVA result show that there is an obvious significant both of forward

and backward analysis. According to Fig. 3, a new factor becomes relevant after a certain time, which indicates that another variability source has appeared, that is Troxeutin-BSA complex. At last, no additional factor is obtained. In the backward, similar information could be obtained from Fig. 3. According to the discussion above, it is reasonable to propose that at least three chemical species active (BSA, troxeutin and Troxeutin-BSA complex) involved in the process. Moreover, it is further confirmed that the fluorescence quenching of BSA caused by troxeutin through a static quenching procedure. The behavior of each chemical species can be obtained by the soft of MCR-ALS (Fig. 4 and 5). These results are good agreement with the previous work, which reported the binding of troxeutin to BSA by absorption and fluorescence spectroscopy. Contrary to previous work, this paper presents the methods combination of synchronous fluorescence spectroscopy and MCR-ALS to study the interaction of troxeutin with BSA and provide more information for understanding their binding characteristics.

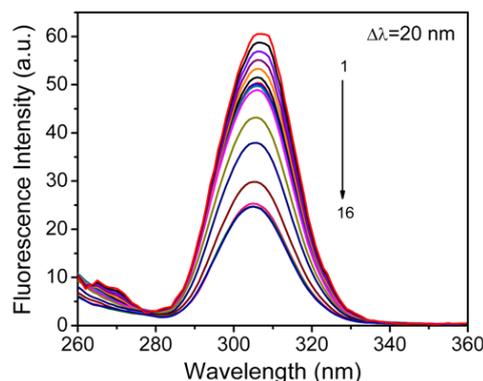


Fig. 1. The synchronous fluorescence spectroscopy of the binding of BSA to troxeutin (the curves from 1 to 16 [Troxeutin]: [BSA] = 0,0.5,1.0,...,7.5)

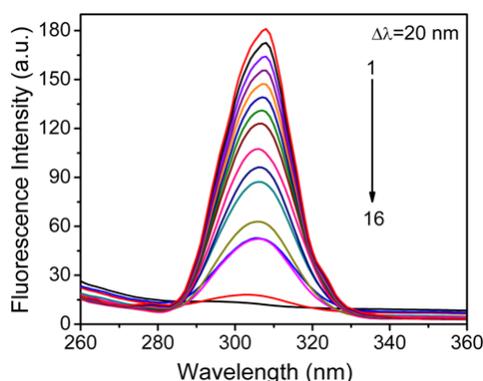


Fig. 2. The synchronous fluorescence spectroscopy of the binding of BSA to troxeutin (the curves from 1 to 16 [BSA]:[Troxeutin] = 7.5, 6.0, 5.5,...,0)

The MCR-ALS Analysis

As the synchronous fluorescence spectra overlap each other (see from Fig. 1 and 2), it is very difficult to determine the possibility of the interaction between troxerutin and BSA. In order to solve this problem, the extended matrix of the synchronous fluorescence spectra data was set up and was analyzed by soft MCR-ALS. Moreover, by using this method, we can obtain the pure substance spectra of the various components based on the MCR-ALS analysis. These results help us to understand the binding mechanism of troxerutin to BSA.

Figure 4 and 5 show the pure substance spectroscopy and the curve of concentration trend, respectively. According Fig. 4 and 5, it can be concluded that there are three chemical species in the Troxerutin-BSA system, which is fully consistent with the results of EFA analysis. The results confirm that troxerutin can interact with BSA from a Troxerutin-BSA complex, thus, it can

be concluded that the quenching mechanism of BSA caused by troxerutin is a static quenching.

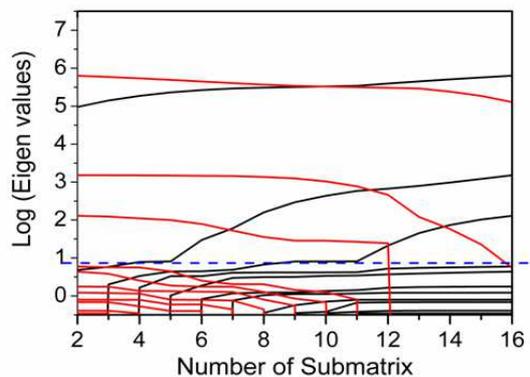


Fig. 3. The EFA plots of the synchronous fluorescence spectroscopy of the binding of BSA to troxerutin. Forward (black lines) and backward (red lines) directions

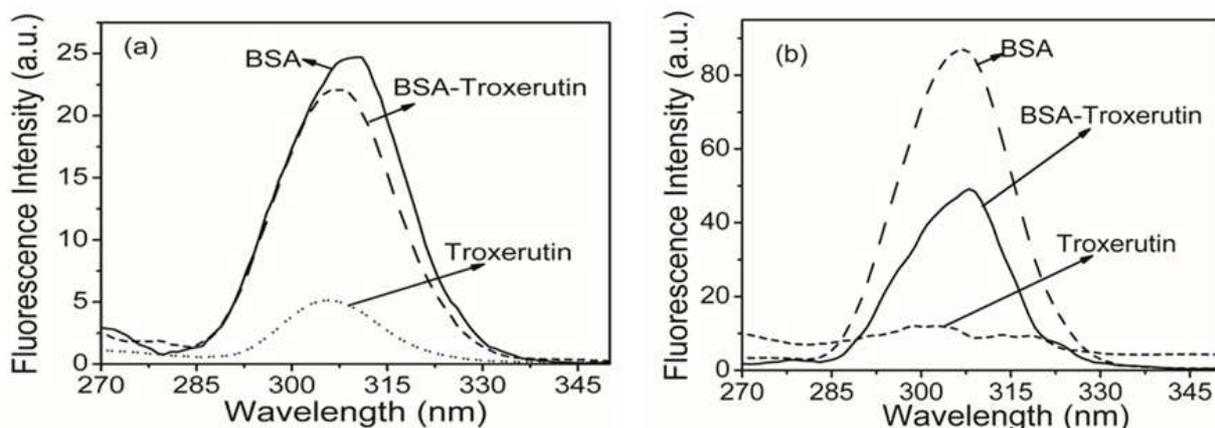


Fig. 4. The pure substance spectroscopy of MCR-ALS analysis (a) experiment 1; (b) experiment 2

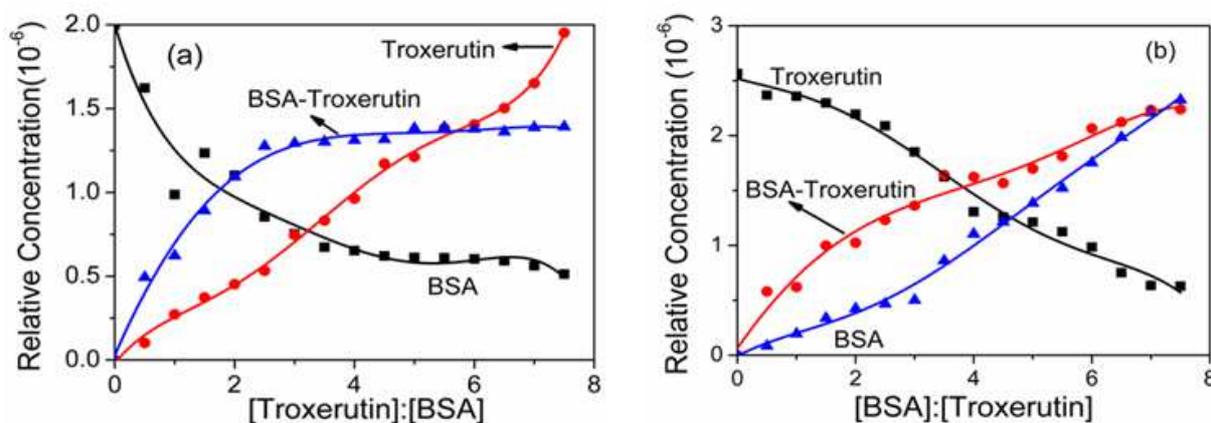


Fig. 5. The concentration trend of MCR-ALS analysis of the interaction between BSA with troxerutin (a) experiment 1; (b) experiment 2

According to Fig. 5a, it can be seen that the concentration of BSA gradually decrease and the complex of Troxerutin-BSA increase with the increasing concentration of troxerutin. From Fig. 5b, the concentration of Troxerutin-BSA complex gradually increase and the degradation of free troxerutin with the added BSA.

Moreover, the concentrations of Troxerutin-BSA complex remained stable as the [BSA]: [Troxerutin] or [Troxerutin]: [BSA] mole ratio in the complex is about 7 and 5, respectively, showing that one troxerutin molecule was bound to one base pair (Anbazhagan and Renganathan, 2008). The average value for the binding constant was obtained from Equation 4:

$$K_{app} = \frac{[BSA - Troxerutin]}{[BSA][Troxerutin]} \quad (4)$$

where, [BSA-Troxerutin], [BSA] and [Troxerutin] is the concentration of free troxerutin-BSA complex, BSA and troxerutin respectively.

According to Equation 4, the average value of the binding constants were calculated as 6.43×10^5 at the [BSA]:[Troxerutin] = 7 and $6.89 \times 10^5 \text{ mol}^{-1} \text{ L}$ at the [Troxerutin]:[BSA] = 5. The prediction of two kinds of dropping models is good agreement with experimental results drawn from MCR-ALS, which indicate that the spectra and concentration profiles of Troxerutin-BSA complex were correctly clarified.

Conclusion

The binding of troxerutin to BSA was studied by combination of the synchronous fluorescence spectroscopy and MCR-ALS method. The EFA and MCR-ALS analysis results showed that there are three chemical species in the Troxerutin-BSA system, detailed analysis revealed that the fluorescence quenching of BSA caused was a static quenching mechanism. Meanwhile, the average binding constant was obtained as $6.43 \times 10^5 \text{ mol}^{-1} \text{ L}$ at the [BSA]:[Troxerutin] = 7 and $6.89 \times 10^5 \text{ mol}^{-1} \text{ L}$ at the [Troxerutin]:[BSA] = 5.

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Author's Contributions

Tianhu Wang: Planning and performing the experiments and data interpretation.

Yi Yan: Data test.

Yinsheng Luo: Data analysis.

Ethics

This article is original and contains unpublished material. The corresponding author confirms that all the other authors have read and approved the manuscript and no ethical issues involved.

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