



Determination of Human Serum Albumin with Tetra-amino Copper Phthalocyanine by Resonance Light Scattering Technique

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Resonance light scattering has been extensively used in the detection of concentrations of biological macromolecules. For the development of resonance light scattering based protein detection sensor at the point-of-care, it is necessary to make the enhanced resonance light scattering wavelength move to long wave direction to improve the detection sensitivity. A method for determination of protein was developed in presence of tetra-amino copper phthalocyanine (TACuPc) by resonance light scattering technique. In acidic solution, the interaction between TACuPc and human serum albumin yielded strongly enhanced resonance light scattering signals at wavelength 454 nm and the enhanced intensity of resonance light scattering was proportional to the concentration of proteins. The effects of pH and interfering species on the determination of protein were examined. Under optimal conditions, the linear range of human serum albumin is 0-2 $\mu\text{g mL}^{-1}$ and the detection limit is 19.37 ng mL^{-1} .

Key Words: Protein, Human serum albumin, Resonance light scattering, Tetra-amino copper phthalocyanine.

INTRODUCTION

The quantitative analysis of protein is of great importance in biochemistry and clinical application because it can provide information for diagnosis of diseases and measure of other components. The traditional methods for the quantitative determination of protein are the Lowry¹, Bradford^{2,3}, bromophenol blue^{4,5} and bromocresol green⁶. Some new methods were also developed, such as spectrophotometric⁷, fluorimetric^{8,9} and chemiluminescent¹⁰ methods. However, most of these methods suffer from the disadvantage of low sensitivity, strong sorption, poor selectivity, the toxicity and high costs of the fluorescence reagents, relatively long analytical time.

Resonance light scattering is a special elastic scattering produced when the wavelength of Rayleigh scattering is located at or close to its molecular absorption. Pasternack *et al.*¹¹ firstly established a sensitive and simple resonance light scattering technique to study the biological macromolecules with a common fluorescence spectrometer. Huang *et al.*¹² employed resonance light scattering technique to establish a high sensitive method for DNA determination. Hence, resonance light scattering technique has become a new interesting

method for the determination of biological molecules, such as nucleic acids^{13,14}, proteins¹⁵, aggregated porphyrin¹⁶ and sugar¹⁷. Commonly, the peak spectral sensitivity of silicon photodetector is located at near infrared. If the protein can enhance the resonance light scattering signal and shift to near infrared, it will create condition for determination of protein with detection sensors at the point-of-care. Phthalocyanines have strong absorption of visible radiation between 600 and 850 nm. In this study, it was found that the tetra-amino copper phthalocyanine (TACuPc) and human serum albumin (HSA) yielded strongly enhanced resonance light scattering signals in an acidic solution. The enhanced intensity of resonance light scattering was proportional to the concentration of human serum albumin. The method has been applied to the determination of total proteins in human serum albumin samples with satisfactory results.

EXPERIMENTAL

Intensity and spectra of resonance light scattering were obtained with an F-2500 spectrofluorimeter (Hitachi Ltd., Tokyo Japan). A quartz cuvette (1 × 1 cm cross-section) was used. The absorption spectra were made on a UV-2300 spectrophoto-

meter (Tianmei, Shanghai, China). All the pH measurements were made with a model pHs-301 pH meter (Xiamen, China).

Tetra-amino copper phthalocyanine was synthesized and purified as described¹⁸. The product was identified by polyamide thin layer chromatography, UV-VIS absorption spectrum, fluorescence spectra and infrared spectrum. A stock solution of 1.0×10^{-4} mol L⁻¹ was prepared in dimethyl sulfoxide.

The standard solutions of human serum albumin (Hua-Mei Biochemical Reagent, China) was directly dissolved in water to prepare stock solutions and stored at 0-4 °C.

Five kinds of buffers, citric acid-sodium citrate, KCl-HCl, Na₂HPO₄-citric acid, potassium biphthalate-HCl and Britton-Robinson [(H₃PO₄, HOAc, H₃BO₃)-NaOH] were used.

All chemical used were of analytical grade and redistilled deionized water was used throughout.

To a 10 mL calibrated tube was added 1 mL of Britton-Robinson buffer solution and an appropriate volume of TACuPc and amount of human serum albumin or sample solution successively, then diluted with water to the mark and mixed thoroughly. The resonance light scattering spectrum was recorded by scanning simultaneously the excitation and emission monochromators ($\Delta\lambda = 0$ nm) of the F-2500 fluorescence spectrophotometer (with 5 nm slit width) from 300 to 700 nm. The resonance light scattering intensity *I* for the reaction product and *I*₀ for the reagent blank were measured at the maximum scattered wavelength, $\Delta I = I - I_0$.

RESULTS AND DISCUSSION

Molecular structure of TACuPc: Fig. 1 is the molecular structure of TACuPc. The structure of its bulk moiety is similar to that of porphyrin, with four benzene rings substituted by a strongly polar amino group, respectively, in the periphery of the bulk moiety. Because of the introduction of four strongly polar amino groups, the solubility of phthalocyanine in water was greatly enhanced.

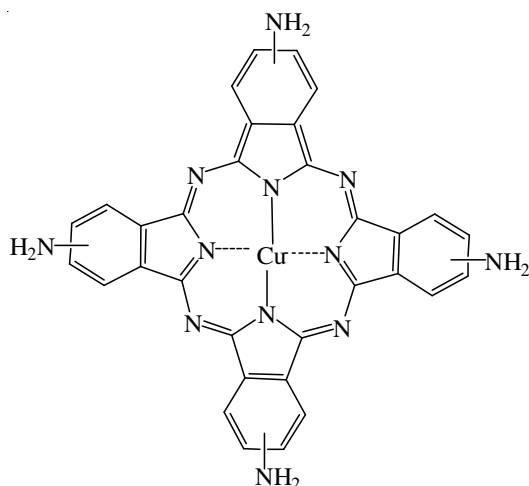


Fig. 1. Molecular structure of TACuPc

Spectral characteristics: The resonance light scattering spectra of TACuPc, human serum albumin and the mixture of TACuPc with human serum albumin are shown in Fig. 2. It can be seen that the resonance light scattering intensities of TACuPc and human serum albumin are quite weak, respectively.

However, when adding human serum albumin to TACuPc or vice versa, a strong resonance light scattering signal can be observed at 454 nm, which indicates that TACuPc and human serum albumin interact with each other. Moreover, the enhanced resonance light scattering intensity increases with the increasing HAS concentration. Therefore, 454 nm was selected as the analytical wavelength.

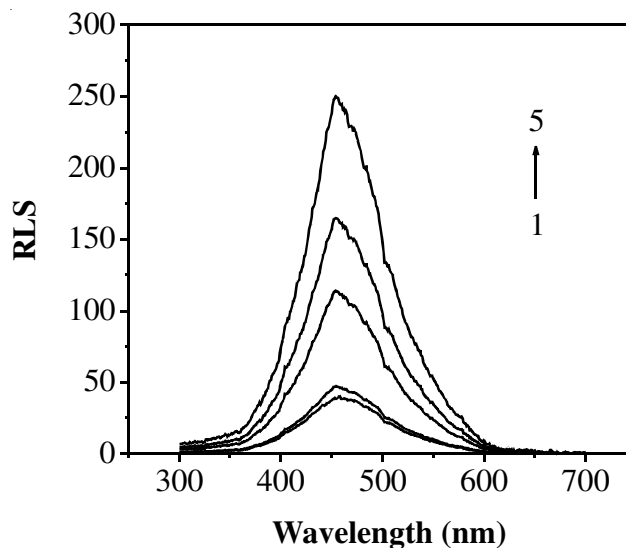


Fig. 2. Resonance light scattering (RLS) of human serum albumin (1); the effect of concentration of human serum albumin on the RLS spectra (2-5). TACuPc: 5.0×10^{-5} mol L⁻¹, human serum albumin ($\mu\text{g mL}^{-1}$): 1, 1.0; 2, 0.0; 3, 0.25; 4, 0.5; 5, 1.0. pH 4.0

Fig. 3 is the absorption spectra of TACuPc in the presence of human serum albumin. It is obvious that TACuPc has two absorption bands, a Soret band and a Q band. As different concentrations of human serum albumin were mixed with TACuPc, no significant wavelength shift occurred for the peaks, however, the absorption peak at 720 nm obviously decreased with increasing amounts of human serum albumin. This further confirmed the strong interaction between TACuPc and human serum albumin.

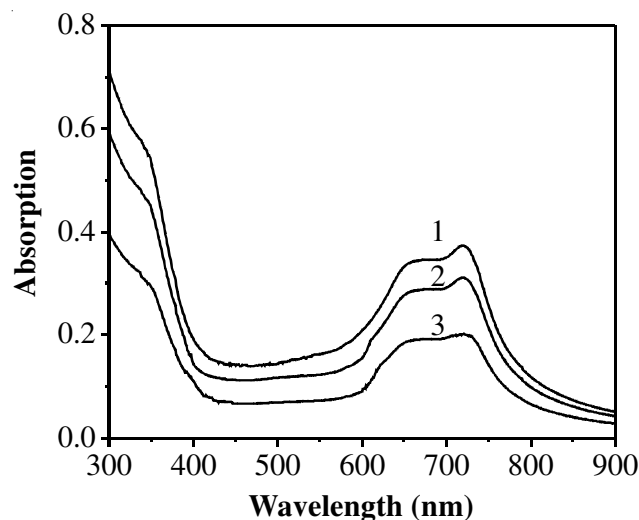


Fig. 3. Absorption spectra of TACuPc in the presence of human serum albumin. TACuPc, 1.0×10^{-5} mol L⁻¹; human serum albumin ($\mu\text{g mL}^{-1}$): 1, 0; 2, 1.0; 3, 2.0

Stability: At room temperature, the intensity of resonance light scattering reaches quickly a maximum when all reagents are mixed together and the scattering intensity is stable for at least 3.0 h. Therefore, 5 min of incubation time was recommended.

Effect of adding sequence of the reagents: The effect of the following adding sequence of the reagents on the sensitivity of this method was studied: buffer + human serum albumin + TACuPc, buffer + TACuPc + human serum albumin and TACuPc + human serum albumin + buffer. The results show the sequence of buffer + human serum albumin + TACuPc gives the best stabilities and intensities of resonance light scattering signals. Mixing the buffer and protein first provides positively charged protein for this combination and improves determination sensitivity. It is easy for negative charged TACuPc in weakly acidic solution (pH 4) to bind positively charged human serum albumin (pH 4.7). Under the neutral condition, if TACuPc is first mixed with human serum albumin then it is difficult for the negatively charged dye to bind the negatively charged human serum albumin.

Effect of pH and buffer: The influence of pH on resonance light scattering intensity was examined in the pH range 1.0-6.5 (Fig. 4). It can be seen that the optimum range of 2-4, an increase in pH leads to a reduction of the resonance light scattering signal. This is possibly related to the isoelectric point of the protein (human serum albumin, pH 4.7). Before the

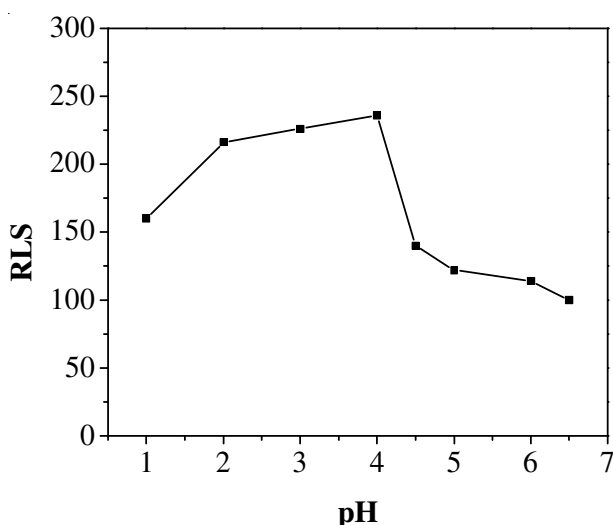


Fig. 4. Effect of pH on resonance light scattering (RLS) intensity. TACuPc, 5.0×10^{-5} mol L⁻¹; human serum albumin, $1.0 \mu\text{g mL}^{-1}$

isoelectric points, protein molecules are in positive charges and react with TACuPc anions by electrostatic interaction. As the pH approaches or exceeds the isoelectric point, protein molecules are neutral or in negative charge, thus the capability to reach with TACuPc is greatly weakened. Therefore, pH 4 was chosen to run the assay.

Experimental results demonstrate that different kinds of buffers, citric acid-sodium citrate, KCl-HCl, Na₂HPO₄-citric acid, potassium biphthalate-HCl and Britton-Robinson, were tested and the results showed a similar changing trend with slightly different effect on the resonance light scattering signal. Britton-Robinson buffer was finally chosen.

Optimum amounts of TACuPc: The effect of TACuPc concentration on scattering intensity was examined. The results are shown in Fig. 5. It can be seen that more sensitivity and the widest linear range are reached at a TACuPc concentration of 5.0×10^{-5} mol L⁻¹. The concentration of 5.0×10^{-5} mol L⁻¹ was recommended.

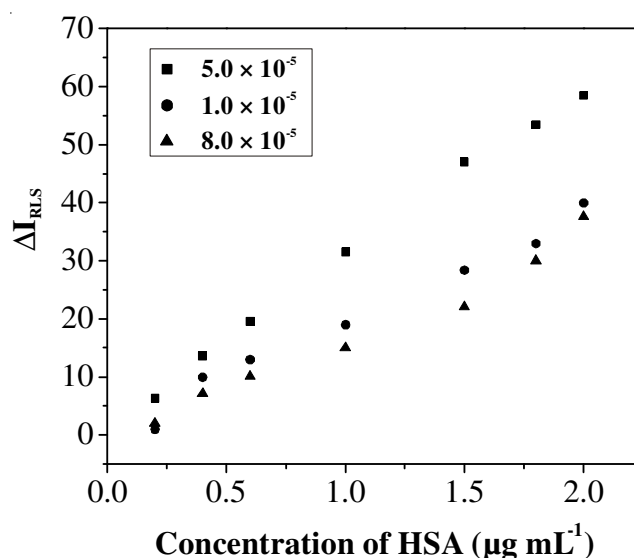


Fig. 5. Effect of TACuPc concentration on the intensity of resonance light scattering (RLS) at pH 4.0, TACuPc ($\times 10^{-5}$ mol L⁻¹)

Interference: The influence of coexisting substance, such as metal ions and amino acids, was tested at $1.0 \mu\text{g mL}^{-1}$ human serum albumin. The experimental results are listed in Table-1. It can be seen that amino acids and ions hardly interfere with the determination.

TABLE-1
RATE CONSTANT (k) FOR $2\text{HI}(\text{g}) \rightarrow \text{H}_2(\text{g}) + \text{I}_2(\text{g})$ AT ABSOLUTE TEMPERATURE (T)^a

Foreign substance	Maximum concentration ($\mu\text{g mL}^{-1}$)	Relative error caused (%)	Foreign substance	Maximum concentration ($\mu\text{g mL}^{-1}$)	Relative error caused (%)
Ca ²⁺ (chloride)	20	2.6	Urea	100	-1.6
K ⁺ (chloride)	25	-2.1	EDTA	20	4.2
Co ²⁺ (chloride)	20	3.9	Glucose	50	-4.7
Fe ³⁺ (chloride)	0.2	4.7	L-Tryptophan	4	3.7
Cd ²⁺ (chloride)	10	1.3	Glycine	10	-2.9
Mg ²⁺ (chloride)	15	2.6	L-Lyrosine	75	4.8
Mn ²⁺ (chloride)	20	1.2	D-Phenylalanine	100	1.5
Al ³⁺ (chloride)	5	1.8	Pb ²⁺ (Nitrate)	5	2.5
Zn ²⁺ (chloride)	1	2.2	L-Arginine	100	4.1
Na ⁺ (chloride)	20	2.4	—	—	—

Effect of ionic strength: In a general system, the enhancement of the resonance light scattering signal strongly depends on the ionic strength solution. The interaction between human serum albumin and TACuPc is mainly a result of electrostatic forces, so the ionic strength should have some effect on this reaction. It was found that the resonance light scattering intensities are stable when the ionic strength is less than 0.1 mol L⁻¹ NaCl but decrease when the ionic strength is more than 0.1 mol L⁻¹. This phenomenon is possibly due to the decrease of the electrostatic forces between TACuPc and human serum albumin, because the shielding effect of the charges on proteins increases with increasing ionic strength, which unprofitable for the formation of the complex.

Calibration graphs: Calibration graphs for the determination of proteins were constructed. Under the above optimal conditions, the difference of the light scattering intensity in the absence and presence of human serum albumin (ΔI) was proportional to the concentration of proteins with a good linear relationship. The linear regression equation is:

$$\Delta I = 1.639 + 29.91C, r = 0.9994$$

A linear range is 0-2.0 $\mu\text{g mL}^{-1}$, the detection limit is 19.37 ng mL⁻¹.

The limit of detection (LOD) was given by the equation $\text{LOD} = \text{KS}_0/S$, where K is a numerical factor chosen according to the confidence level desired, S_0 is the standard deviation of the blank measurements ($n = 9$, $K = 3$) and S is the slope of the calibration curve. The relative standard deviations (RSD) for 9 determinations of 1 $\mu\text{g mL}^{-1}$ human serum albumin is 2.1 %.

Determinations of total protein in human serum samples: Under the experimental conditions, the present method was used to determine total proteins in human serum samples obtained from the North University of China Hospital, stored at 0-4 °C and diluted 1000-fold with deionized water for analysis. The results for the human serum samples are shown in Table-2. It can be seen that the results obtained by clinical physicians, which indicates this method has the potential in clinical assay.

TABLE-2
DETERMINATION OF PROTEIN IN HUMAN SERUM^a

Sample	Present method (mg mL ⁻¹)	RSD	Clinical data (mg mL ⁻¹)
1	81.3	2.1	81.6
2	76.8	2.3	76.9
3	88.1	3.4	87.7
4	82.4	3.9	83.1

^aEach result was the average of five measurement

Conclusion

In the acidic environment, human serum albumin and negatively organic dye can produce electrostatic forces and form organic dye-protein complexes. At pH 4.0, the interaction between human serum albumin and TACuPc yields strongly enhanced resonance light scattering signals. This new method using an ordinary spectrofluorimeter is sensitive and simple. It is shown that this method has great potential, high sensitivity, good stability, increased simplicity and lower detection limits for the determination of protein. And the method will create conditions for detection sensors at the point-of-care with silicon photodetectors.

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