

## The Effect of Caffeine on the interaction between Bovine Serum Albumin and Losartan potassium by Spectroscopic Methods

Zeinab M. Anwer<sup>1</sup>, Gasser M. Kairy<sup>1</sup>, Hend M. Moustafa<sup>2</sup>, and Nada A. Salman<sup>3</sup>

<sup>1</sup> Department of Chemistry, Faculty of Science, Suez Canal University, Egypt.

<sup>2</sup> Department of Chemistry, Faculty of Science, El-Arish University, Egypt.

<sup>3</sup> Department of Basic Science, Faculty of Dentistry, Sinai University, Egypt.

### ARTICLE INFO

#### Article history:

Received 22 March 2024

Received in revised form 13 April 2024

Accepted 20 April 2024

Available online 20 May 2024

#### Keywords

Absorption,  
Fluorescence,  
Bovine serum albumin,  
Losartan potassium,  
caffeine,  
competitive binding

### ABSTRACT

Losartan potassium (LP) is an antihypertensive drug, and its binding to plasma proteins determines its pharmacokinetic properties. An exogenous substance like Caffeine (CAF) may affect the binding of drugs to proteins. Accordingly, we aim to characterize LP and bovine serum albumin (BSA) interactions in CAF presence employing spectroscopic techniques (UV-vis absorption and fluorescence measurements) under physiological circumstances. Our findings indicated that LP's quenching effect on BSA decreased by adding CAF by calculating the Stern-Volmer constant. Therefore, the binding constants for the BSA-LP and BSA-CAF-LP systems were determined using the Modified Stern-Volmer equation; the results were  $4.47 \times 10^3$  and  $0.23 \times 10^3$ , respectively.

Thermodynamic parameters, the Gibbs free energy ( $\Delta G$ ), enthalpy ( $\Delta H$ ), and entropy ( $\Delta S$ ) calculated for each system show spontaneous interaction for both systems and suggest that the binding interaction between BSA and LP is exothermic and entropy is unfavorable, while for BSA-CAF-LP is endothermic and entropy is favorable.

### 1. Introduction

The drug and protein interactions within the bloodstream significantly affect the pharmacodynamics and pharmacokinetics of therapeutic molecules(1,2). The degree to which a pharmaceutical compound interacts with a protein determines its stability and potential for toxicity(1). About 60% of the total protein content in the human body is made up of albumin, so it is considered the main blood plasma protein(3). Albumin transports unesterified fatty acids, metabolites, organic compounds, dyes, and drugs(4) thereby modulating their delivery, distribution, and efficiency(5). Human serum albumin (HSA) and bovine serum albumin (BSA) share structural similarities (6,7). Both proteins possess three unique drug binding sites: I, II, and III in subdomains IIA, IIIA, and IB, respectively(8–11). In addition to its homogeneity, this substance exhibits other notable attributes, including its exceptional affinity for ligand binding, accessibility, and affordability(12). In our work, we have chosen BSA as a protein model.

Losartan Potassium (LP) is a pharmacological agent utilized to treat hypertension. Its mechanism of action involves the selective and competitive inhibition of type 1 (AT1) angiotensin II receptors, resulting in a reduction of blood pressure(13,14).

LP exhibits a strong affinity for serum albumin, with a binding capacity of around 98.8% (15,16). At a lower drug-to-BSA ratio, LP exhibits a greater affinity for subdomain IIIA in site II of the BSA molecule compared to subdomain IIA in site I. In contrast, when the ratio is increased, it exhibits binding affinity towards both locations (I & II)(15–17). CAF is a natural substance found in the seeds and fruits of more than 63 plant species globally(18,19), coffee(20), cocoa beans(21,22), chocolate(23), cola nuts(24), tea(25), and pharmacologically active substance(26). According to existing literature, it has been established that CAF is the most extensively ingested psychoactive compound on a global scale(27,28). CAF has been revealed to exhibit binding affinity towards site I, corresponding to subdomain IIA of BSA(29–32). As with any exogenous substances, it can impact the pharmacokinetic characteristics of medicines through their reversible binding to serum albumin(33,34). The present paper aims to explore the potential interaction of caffeine with bovine serum albumin (BSA) in the presence of losartan potassium through spectroscopic techniques (UV-visible and fluorescence measurements), to find if there is any hindrance effect of caffeine on the binding of the antihypertensive drug with BSA. There are no previous data on the effect of caffeine on the binding of LP with BSA.

\* Corresponding author at Sinai University

E-mail addresses [nada.ali@su.edu.eg](mailto:nada.ali@su.edu.eg) (Nada A. Salman)

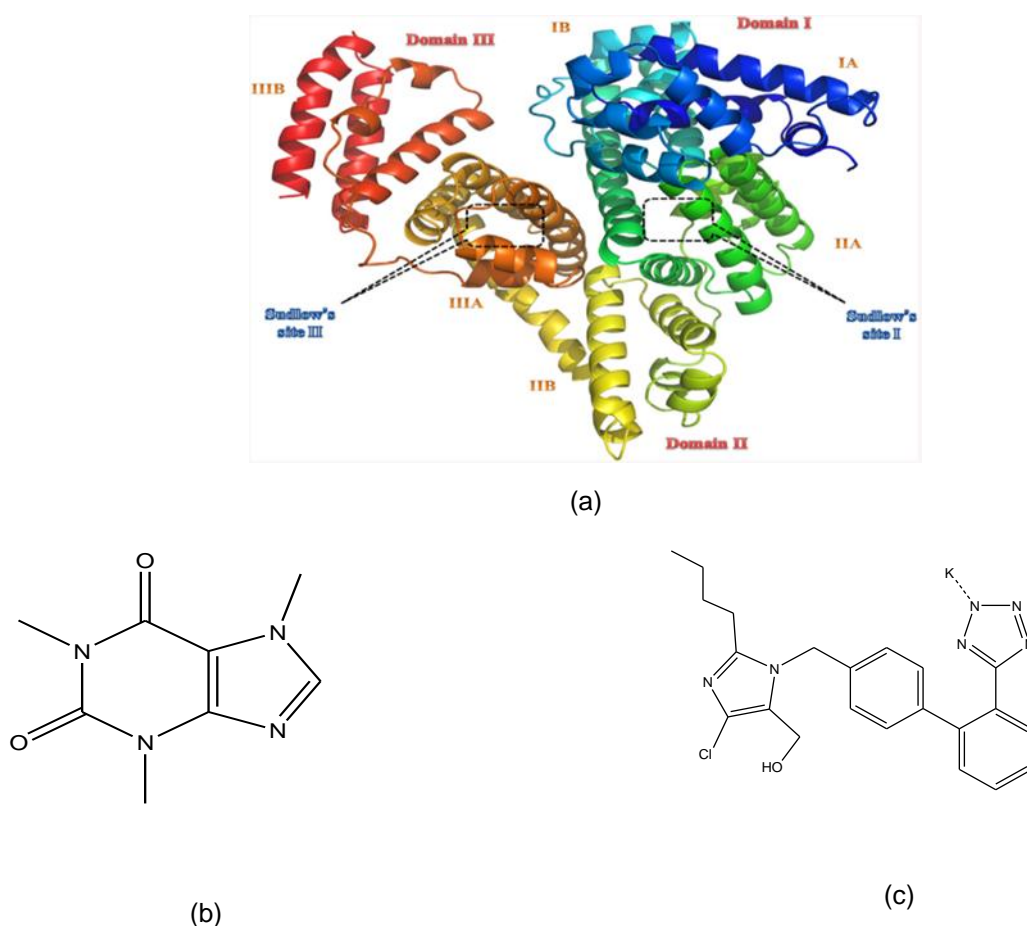
## 2. Materials and methods

### 2.1. Chemicals and reagents

Our study obtained lyophilized BSA powder from Sigma Aldrich ([www.sigmaaldrich.com](http://www.sigmaaldrich.com)) with a purity of at least 96%, ensuring it was free from any fatty acids. The LP sample was received as a complimentary sample from Mepaco-Pharma ([www.mepaco-pharma.net](http://www.mepaco-pharma.net)). The CAF compound was purchased from Sisco Research Laboratories ([www.srlchem.com](http://www.srlchem.com)). The structures of these compounds are provided in Figure 1. Tris-HCl buffer was obtained from Sigma Aldrich ([www.sigmaaldrich.com](http://www.sigmaaldrich.com)).

### 2.2. Sample preparation

Herein, we dissolved 70 mg of BSA in a Tris-HCl buffer (0.1 M, pH 7.4) to prepare a BSA stock solution (0.062 mM), stored at 4°C. The molar concentration of BSA was calculated using a spectrophotometric approach, utilizing the molar extinction coefficient of BSA ( $43,824 \text{ M}^{-1} \text{ cm}^{-1}$ ) at 280 nm(35,36). Stock solutions of (1mM) for each LP and CAF were prepared by dissolving 4.61 and 1.94 mg in Tris-HCl buffer (0.1 M, pH 7.4).



**Fig. 1.** Chemical structure of (a) BSA structure (PDB ID: 3 V03), (b) Caffeine, and (c) Losartan potassium.

### 2.3. Spectroscopic studies

#### 2.3.1. Fluorescence measurements

The fluorescence measurements were carried out using a Jasco FP-6300 spectrofluorometer (<https://jscoinc.com>) with a 150 W Xenon lamp and a quartz cell with a 1.00 cm path length. The aim was to monitor the influence of CAF on the binding of LP with BSA. BSA was kept at a constant concentration of  $1 \mu\text{M}$  and was titrated with various concentrations of LP (0-24

$\mu\text{M}$ ) in Tris-HCl buffer at room temperature. Then a solution consisting equimolar of BSA and CAF was titrated by increasing the LP concentration from 0-24  $\mu\text{M}$  in Tris-HCl buffer at RT. Subsequently, every sample was stimulated at 280 nm, measuring the resulting emission spectra within 290–420 nm. The measurements were conducted using medium response, high sensitivity, and a bandwidth of 5 nm for excitation and emission. Scanning curves were created for each experiment. The data underwent analysis

and processing using Origin Lab 2019 software. Moreover, the temperature effect was observed through repeated steady-state fluorescence measurements at 298, 303, 308, 313, and 318 K. For the BSA-LP system, BSA of 1 μM was incubated with and without various LP concentrations (1, 2, 3, 4, 5, 6, and 7 μM). In contrast, for BSA-CAF-LP interactions, BSA-CAF was incubated alone and with different amounts (1 μM to 7 μM) of LP. Then, the samples were subjected to excitation at 280 nm, followed by recording the emission spectra at 345nm. Stern-Volmer equations were employed for analyzing the data(12,37).

$$\frac{F_0}{F} = 1 + K_{sv}[Q] \tag{1}$$

$$Kq = \frac{K_{sv}}{\tau_0} \tag{2}$$

F<sub>0</sub> and F refer to fluorescence intensity without and with a quencher, respectively; K<sub>sv</sub> is the Stern–Volmer quenching constant; Kq refers to the bimolecular rate constant of the quenching reaction; [Q] is quencher concentration; and τ<sub>0</sub> is the protein average integral fluorescence lifetime without quencher, which is in the order of 10<sup>-8</sup>s(38). The binding constant and the number of binding sites were studied by the modified Stern-Volmer equation (39,40).

$$\text{Log}(F_0-F)/F = \text{Log} K + n \text{log}[Q] \tag{3}$$

F<sub>0</sub> and F refer to fluorescence intensity without and with a quencher, respectively; K is the binding constant; n is the number of binding sites; and [Q] is the quencher concentration.

$$\ln K_b = -\left(\frac{\Delta H}{RT}\right) + \left(\frac{\Delta S}{R}\right) \tag{4}$$

$$\Delta G = \Delta H - T\Delta S = -R\ln K_b \tag{5}$$

ΔH and ΔS are the enthalpy and entropy changes, respectively; K is the binding constant at the corresponding temperature; and R is the gas constant (41).

**2.3.2. UV-Vis studies**

Ultraviolet-visible (UV-vis) absorption spectroscopy is used for understanding protein-ligand interactions(42).

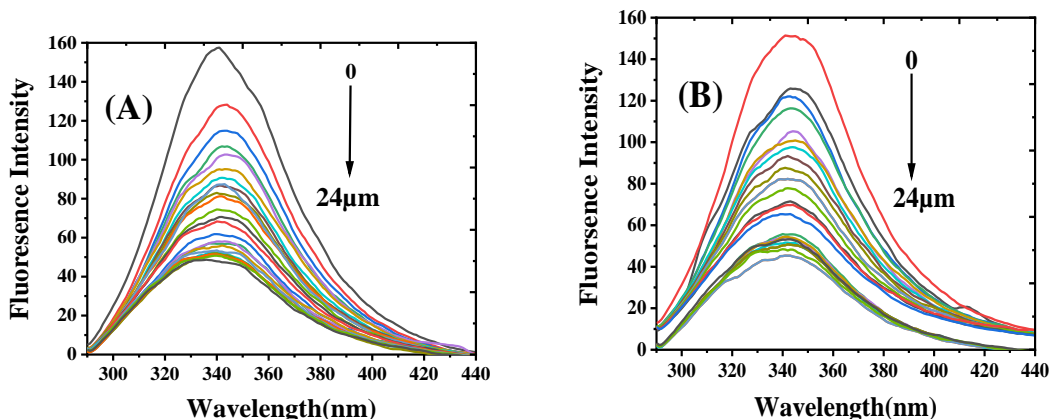
Conformational changes of the protein can be checked through UV-vis spectral examinations. UV-1800 Shimadzu double-beam UV-Vis spectrophotometer with a 1.00 cm quartz cell (<https://www.shimadzu.com>) was employed to measure the absorption spectra. BSA (0.01mM) was titrated by LP (0.01mM) and followed by the same concentration of CAF. The absorption of free CAF and LP are also taken into account. The samples were prepared in a Tris-HCl buffer with a pH of 7.4. Measurements were estimated within the wavelength range of 220-320.

**3. Results and Discussion**

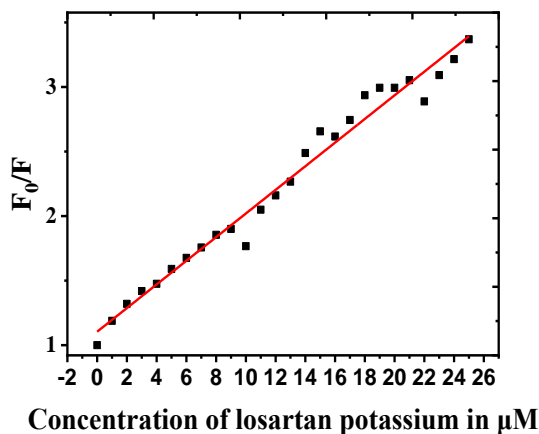
**3.1. Fluorescent measurements**

**3.1.1 Binding of BSA with LP in the absence and presence of CAF**

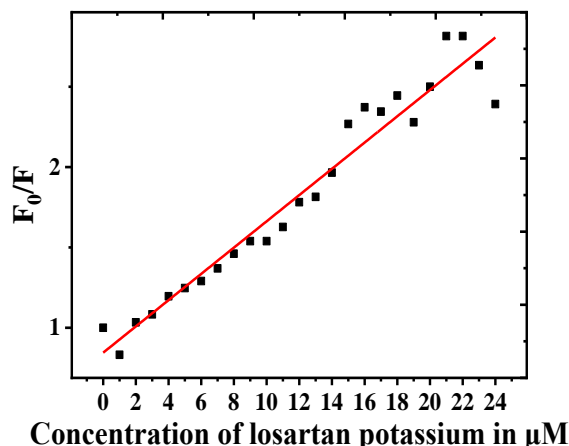
Previous studies have demonstrated the binding of both LP and CAF to BSA in subdomain IIA at site I. Therefore, our objective was to investigate any possible influence of CAF on LP-BSA binding. LP was found to reduce the fluorescence of BSA when CAF is not present with a quenching constant value of 12.7 × 10<sup>4</sup> mol<sup>-1</sup>L at 298 K. On the other hand, when CAF is present the quenching constant value was 6.27 × 10<sup>4</sup> mol<sup>-1</sup>L at 298 K (Figure 2.3). The decrease in the quenching ability of LP, as shown by the significant decrease in the Ksv value, when CAF is present compared to BSA-LP alone, suggests that CAF and LP compete for binding at site I in domain IIA. As a result, the interaction of LP with BSA in the presence of caffeine is reduced, likely due to the probable binding of caffeine with losartan potassium. This greatly decreases the binding of the LP drug with BSA, especially at room temperature. To conduct a more comprehensive analysis of the impact of CAF on the interaction between BSA and LP, the binding constant values were determined by employing a modified Stern-Volmer equation. The binding constant was found to be 4.47×10<sup>3</sup> for BSA-LP and 0.23×10<sup>3</sup> for BSA-CAF-LP Figure 4, suggesting that CAF hinders the binding of LP to BSA.



**Fig. 2.** Emission spectra of (A) 1μM BSA in the presence of various concentrations of Losartan Potassium and (B) BSA-CAF system through various Losartan Potassium concentrations in Tris-HCl buffer=7.4.

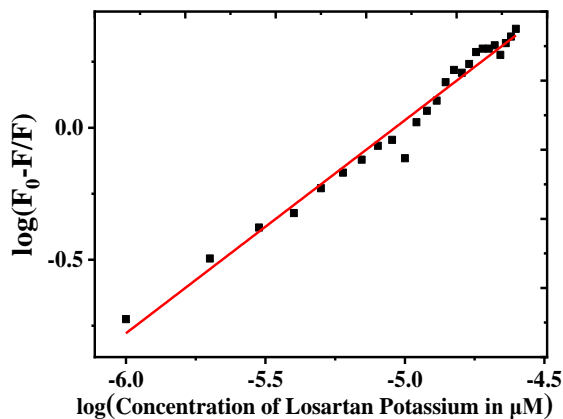


(A)

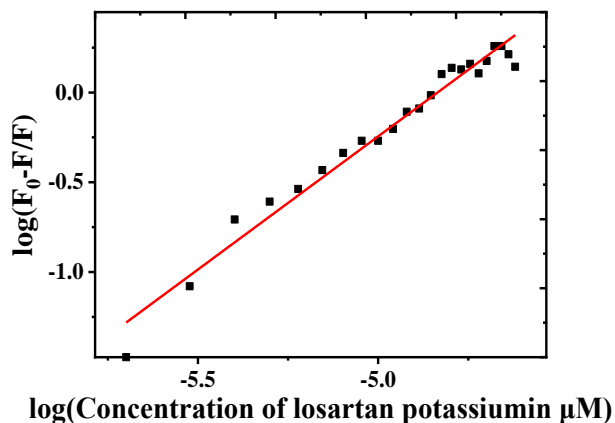


(B)

**Fig. 3.** Stern Volmer plot for (A) the binding of LP with BSA in Tris-HCl buffer=7.4. (B) the binding of BSA with losartan potassium in the presence of caffeine in Tris-HCl buffer=7.4.



(A)



(B)

**Fig. 4.** Modified Stern Volmer plot for (A) the binding of LP with BSA Tris-HCl buffer pH=7.4 and (B) binding of BSA with losartan potassium in the presence of caffeine in Tris-HCl buffer=7.4.

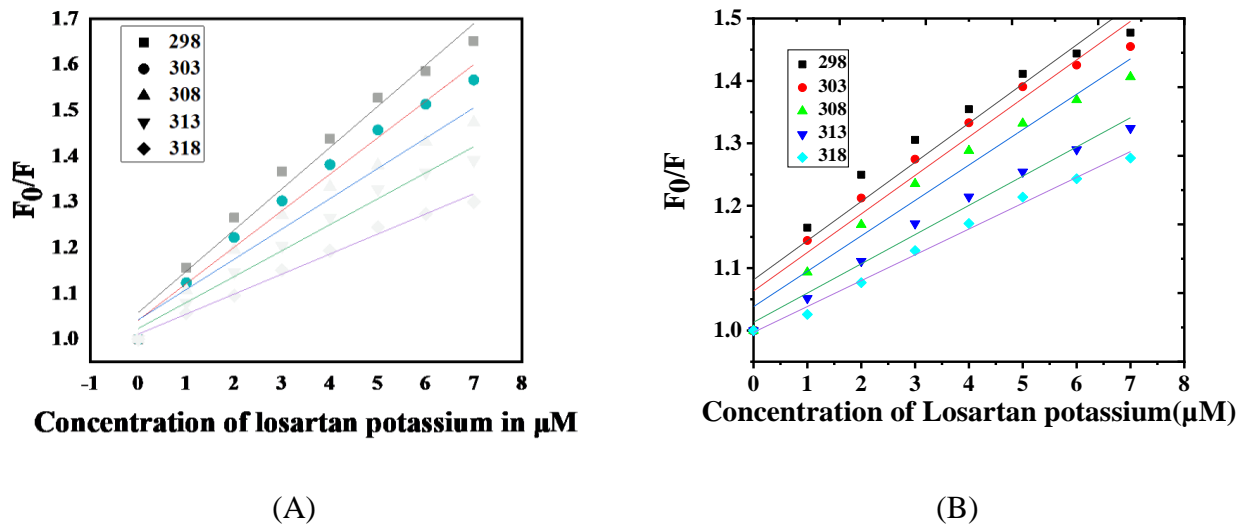
### 3.1.2 Determination of the quenching mechanism of BSA-LP and BSA-CAF-LP system

The decrease in fluorescence intensity results from various molecular interactions, including the formation of ground state complexes, rearrangement of excited state molecules, and energy transfer. Dynamic quenching, closely tied to diffusion, occurs due to these interactions. Higher temperatures accelerate diffusion, resulting in elevated quenching constants. Conversely, in static quenching, increased temperatures prompt the breakdown of complexes formed between drugs and proteins, consequently reducing quenching constant values. (43).

Fluorescence spectroscopy was performed at 298 K, 303 K, 308 K, and 313 K to investigate the quenching mechanism related to the interaction between LP and BSA, as well as LP with BSA-CAF based on their observed fluorescence quenching. The linearity of the Stern-Volmer plots for the BSA-LP and BSA-CAF-LP systems at five different temperatures can be shown in Figure 5, namely in the relationship between  $F_0/F$  and LP concentration [M]. The obtained  $K_{sv}$  values exhibited an inverse correlation with temperature for both systems (Table 1) revealing that quenching is static. Additionally, a graphical representation was generated to illustrate the relationship between the logarithm of the expression  $[(F_0-F)/F]$  and the logarithm of

LP concentration [M]. Subsequently, the intercepts of these graphs were employed to calculate the apparent binding constant (K) for the BSA-LP and BSA-CAF-LP systems across five different temperatures. Figure 6 and Table 2 indicate a significant decrease as temperature increases in the case of the BSA-LP system, indicating that a

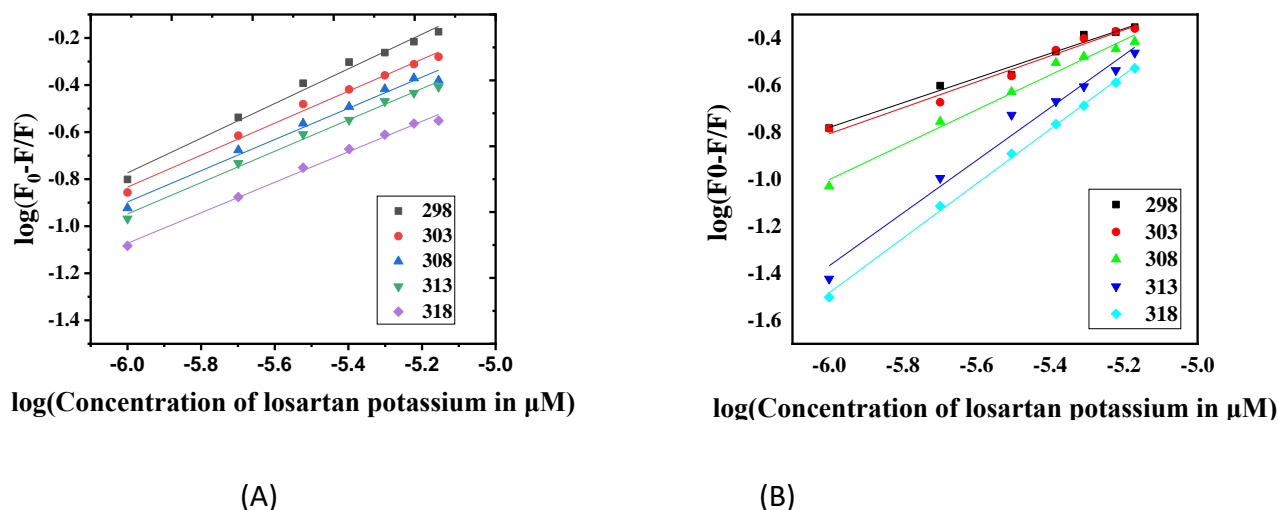
temperature rise disrupts the complex formation between BSA and LP. While K values exhibited notable increases with higher temperatures for the interaction of LP with BSA-CAF, implying that the interaction becomes stronger under elevated temperature conditions.



**Fig. 5.** Stern–Volmer plot for (A) LP with BSA binding and (B) BSA binding with losartan potassium in caffeine presence at different temperatures.

**Table 1** Stern–Volmer Quenching constant for BSA-LP and BSA-CAF-LP at different temperatures.

System	Temperature (K)	Ksv×10 <sup>4</sup> (mol <sup>-1</sup> L)	R <sup>2</sup>
BSA-LP	298 ° K	12.7	0.968
	303 ° K	12.3	0.972
	308° K	11.7	0.974
	313 ° K	11.2	0.979
	318 ° K	10.6	0.988
BSA-CAF-LP	298 ° K	6.27	0.914
	303 ° K	6.17	0.942
	308° K	5.67	0.966
	313 ° K	4.68	0.986
	318 ° K	4.13	0.991



**Fig. 6.** Modified Stern–Volmer plot for (A) LP with BSA binding and (B) BSA binding with losartan potassium in caffeine presence in Tris-HCl buffer=7.4 at different temperatures.

**Table 2** Binding constant of BSA-LP and BSA-CAF-LP at different temperatures.

System	Temperature (K)	$K \times 10^3$ (mol <sup>-1</sup> L)	n	R <sup>2</sup>
BSA-LP	298 ° K	4.47	0.73	0.986
	303 ° K	1.74	0.67	0.990
	308° K	1.21	0.66	0.980
	313 ° K	1.12	0.66	0.991
	318 ° K	0.65	0.64	0.993
BSA-CAF-LP	298 ° K	0.02	0.52	0.976
	303 ° K	0.03	0.55	0.975
	308° K	0.28	0.74	0.981
	313 ° K	22.3	1.11	0.974
	318 ° K	29.6	1.15	0.997

**3.1.3 Determination of thermodynamic parameters and nature of binding forces involved in BSA-LP and BSA-CAF-LP systems**

The thermodynamic parameters (e.g., free energy (G), entropy (S), enthalpy (H)) for the BSA-LP and BSA-CAF-LP systems were studied at five different temperatures: 303K, 308K, 313K, and 318K. A plot (Fig. 7) of the natural logarithm of the binding constant (ln Kb) against the reciprocal of temperature (1/T) demonstrates that ΔG is

consistently negative across all five temperatures for the binding of LP to BSA and BSA-CAF. This negative value suggests that the interaction is spontaneous for both systems. Similarly, ΔH and ΔS values for BSA-LP were -67.84 kJ mol<sup>-1</sup> and -159.76 J K<sup>-1</sup> mol<sup>-1</sup>, respectively. The presence of negative values for both ΔS and ΔH suggests that the binding interaction between BSA and LP is exothermic and driven mainly by enthalpy, as the entropy is unfavorable here showing that the van der Waals forces



and hydrogen bonds are crucial in the BSA-LP interaction. In contrast,  $\Delta H$  and  $\Delta S$  values were  $328.09 \text{ kJ mol}^{-1}$  and  $1139.07 \text{ J K}^{-1} \text{ mol}^{-1}$ , respectively, suggesting that the binding interaction between BSA-CAF and LP is endothermic and driven mainly by enthalpy and the acting force is hydrophobic interaction.

### 3.2. UV-vis spectrophotometric measurements

UV-Vis absorption spectroscopy is a straightforward and widely utilized technology frequently employed for examining structural alterations and investigating complex development(44). The alterations, such as the hypochromic or hyperchromic impact and the red or blue shift in the UV spectra throughout titration, can indicate the prevailing interaction mode between chemicals and BSA. Dynamic

(collisional) quenching exclusively affects the fluorophore-excited states, leaving the absorption spectra unaffected. Nonetheless, developing a ground-state complex leads to modifications in the fluorophore absorption spectrum (45). BSA's absorption peak of nearly 280 nm is attributed to the aromatic amino acid residue tryptophan (46,47). Figure 8 shows the absorption spectra of free CAF and LP at identical concentrations (0.01mM) and their interaction with BSA. The observed increase in absorbance at approximately 280 nm is mainly attributed to the  $\pi-\pi^*$  transition of aromatic amino acid residues present in BSA (48,49) which means that both alter the environment of BSA tryptophan residues.

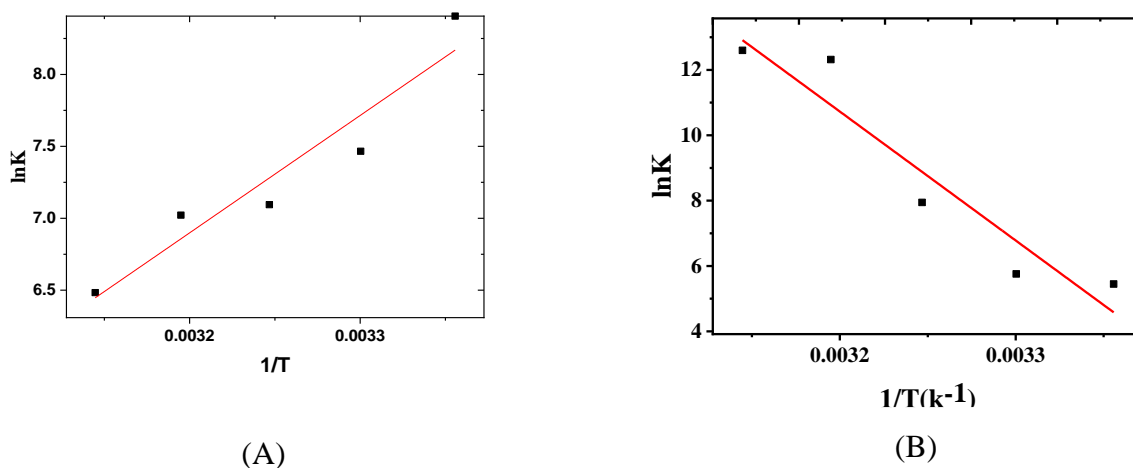


Fig. 7. Van't Hoff equation for binding (A) LP with BSA and (B) LP with BSA-CAF.

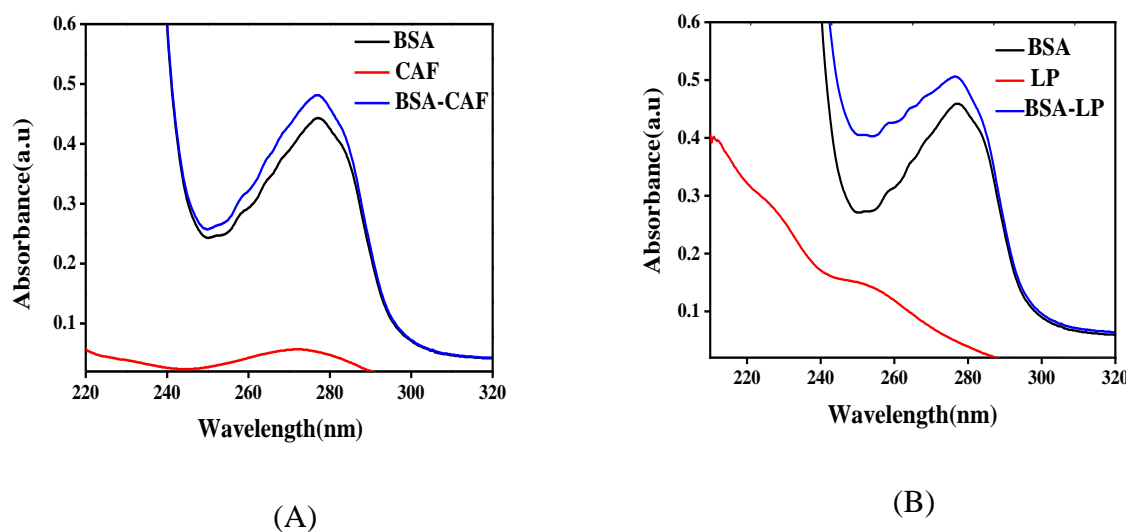
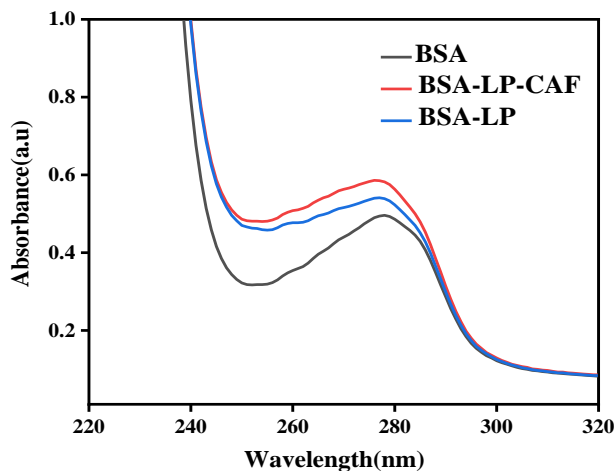


Fig. 8. Absorption spectra of BSA(0.01mM) in (A) caffeine and (B) losartan potassium presence each of Tris-HCl buffer.

The absorption spectra of BSA with LP were accurately measured, and then 0.01 mM of CAF was introduced Figure 9. Our findings indicate that adding LP, CAF, or a combination of both heightens BSA absorbance values. This suggests the formation of stable complexes in all

instances, further supported by a similar observation made using the fluorescence method. The UV-Vis spectral results demonstrate that adding CAF to BSA-LP increases absorbance, providing evidence for developing novel protein-drug-drug complexes (BSA-LP-CAF).



**Fig. 9.** Absorption spectra of 0.01m BSA, (1:1) BSA–LP system of 0.01mM, and (1:1:1) BSA-LP-CAF of 0.01mM in Tris-HCl buffer.

**Table 3** Thermodynamic parameters for interaction of BSA-LP and BSA-CAF-LP complex at different temperatures.

System	Temperature (K)	$\Delta G$ (kJ mol <sup>-1</sup> )	$\Delta H$ (kJ mol <sup>-1</sup> )	$\Delta S$ (J K <sup>-1</sup> mol <sup>-1</sup> )
BSA-LP	298 ° K	-20.24	-67.84	-159.76
	303 ° K	-19.44		
	308° K	-18.64		
	313 ° K	-17.84		
	318 ° K	-17.04		
BSA-CAF-LP	298 ° K	-11.35	328.09	1139.07
	303 ° K	-17.04		
	308° K	-22.74		
	313 ° K	-28.43		
	318 ° K	-34.13		

**Conclusion**

Serum albumins possess binding sites for many medications, with Sudlow sites I and II being the primary pharmaceutical binding sites (11,50). This study aims to investigate the impact of pharmacological medicines,

including CAF, on the binding of LP to BSA, specifically at subdomain IIA, where LP is known to bind. Our findings indicate a significant impact of CAF on LP binding to BSA. The relationship between the proportion of unbound medication and serum albumin has been widely



acknowledged as a determinant of pharmacological response (51). It is not recommended for patients undergoing LP administration to concomitantly consume CAF from any source, including drugs, food, or beverages. This precaution is necessary due to the potential for mutual nullification of therapeutic effects or the occurrence of unfavorable pharmacological responses.

The findings of our study indicate that CAF presence can affect the LP binding constant with BSA, potentially affecting the therapeutic efficacy of the medicine. Consequently, healthcare professionals should caution patients against the concurrent use of CAF and LP during therapy.

### Acknowledgment

There is nothing to declare. The contributing author supported the work.

### Conflicts of Interest

The authors declare that they have no conflicts of interest in this work.

### References

- Rahman Y, Afrin S, Tabish M. Interaction of pirenzepine with bovine serum albumin and effect of  $\beta$ -cyclodextrin on binding: A biophysical and molecular docking approach. *Arch Biochem Biophys* [Internet]. 2018 Aug 15;652:27–37. Available from: <https://doi.org/10.1016/j.abb.2018.06.005>
- Rahman Y, Afrin S, Alhaji Isa M, Ahmed S, Tabish M. Elucidating the molecular interaction of serum albumin with nizatidine and the role of  $\beta$ -cyclodextrin: a multi-spectroscopic and computational approach. *J Biomol Struct Dyn* [Internet]. 2020 Mar 23;38(5):1375–87. Available from: <http://dx.doi.org/10.1080/07391102.2019.1604265>
- Afrin S, Rahman Y, Alhaji Isa M, Ahmed S, Tabish M. Biophysical insights into the binding characteristics of bovine serum albumin with dipyrindamole and the influence of molecular interaction with  $\beta$  cyclodextrin. *J Biomol Struct Dyn* [Internet]. 2020 Jul 2;38(10):3046–58. Available from: <http://dx.doi.org/10.1080/07391102.2019.1651220>
- Varshney A, Sen P, Ahmad E, Rehan M, Subbarao N, Khan RH. Ligand binding strategies of human serum albumin: How can the cargo be utilized? *Chirality* [Internet]. 2010 Jan;22(1):77–87. Available from: <https://onlinelibrary.wiley.com/doi/10.1002/chir.20709>
- Trainor GL. The importance of plasma protein binding in drug discovery. *Expert Opin Drug Discov* [Internet]. 2007 Jan 16;2(1):51–64. Available from: <http://www.tandfonline.com/doi/full/10.1517/17460441.2.1.51>
- Chruszcz M, Mikolajczak K, Mank N, Majorek KA, Porebski PJ, Minor W. Serum albumins-unusual allergens. *Biochim Biophys Acta* [Internet]. 2013 Dec;1830(12):5375–81. Available from: <https://linkinghub.elsevier.com/retrieve/pii/S0304416513002766>
- Carter DC, Ho JX. Structure of Serum Albumin. In: *Advances in Protein Chemistry* [Internet]. 1994. p. 153–203. Available from: <https://linkinghub.elsevier.com/retrieve/pii/S0065323308606403>
- Carter DC, Chang B, Ho JX, Keeling K, Krishnasami Z. Preliminary Crystallographic Studies of Four Crystal Forms of Serum Albumin. *Eur J Biochem* [Internet]. 1994 Dec;226(3):1049–52. Available from: <https://onlinelibrary.wiley.com/doi/10.1111/j.1432-1033.1994.01049.x>
- Huang BX, Kim HY, Dass C. Probing three-dimensional structure of bovine serum albumin by chemical cross-linking and mass spectrometry. *J Am Soc Mass Spectrom*. 2004 Aug 1;15(8):1237–47.
- Ketrat S, Japrun D, Pongprayoon P. Exploring how structural and dynamic properties of bovine and canine serum albumins differ from human serum albumin. *J Mol Graph Model* [Internet]. 2020;98:107601. Available from: <https://doi.org/10.1016/j.jmgm.2020.107601>
- Bujacz A. Structures of bovine, equine, and leporine serum albumin. *Acta Crystallogr Sect D Biol Crystallogr* [Internet]. 2012 Oct 1;68(10):1278–89. Available from: <http://scripts.iucr.org/cgi-bin/paper?S0907444912027047>
- Jiang TY, Zhou KL, Lou YY, Pan D qi, Shi JH. Probing the behavior of bovine serum albumin upon binding to atenolol: insights from spectroscopic and molecular docking approaches. *J Biomol Struct Dyn* [Internet]. 2018 Apr 4;36(5):1095–107. Available from: <http://dx.doi.org/10.1080/07391102.2017.1311805>
- Burnier M, Wuerzner G. Pharmacokinetic evaluation of losartan. *Expert Opin Drug Metab Toxicol* [Internet]. 2011 May 22;7(5):643–9. Available from: <http://www.tandfonline.com/doi/full/10.1517/17425255.2011.570333>
- Goa KL, Wagstaff AJ. Losartan Potassium. *Drugs* [Internet]. 1996 May;51(5):820–45. Available from: <http://link.springer.com/10.2165/00003495-199651050-00008>
- Sica DA, Gehr TWB, Ghosh S. Clinical pharmacokinetics of losartan. *Clin Pharmacokinet*. 2005;44(8):797–814.
- Ferdosi Kabir, A.; Nazim Uddin, K.; Nazmus Sadat, A.F.M.; Mahboob, Hossain; Abdul Mazid M. Interaction of Palmitic Acid with Losartan Potassium at the Binding Sites of Bovine Serum Albumin. *Ars Pharm* [Internet]. 2010;51(1):28–36. Available from: <http://hdl.handle.net/10481/60183>
- Sun M, Su M, Sun H. Journal of Chemical and Pharmaceutical Research, 2015, 7 ( 10 ): 924-932 Research Article Spectroscopic study on the interaction of losartan potassium and bovine serum albumin. 2015;7(10):924–32.
- Mazzafera P, Silvarolla MB. Caffeine content variation in single green Arabica coffee seeds. *Seed Sci Res*. 2010;20(3):163–7.
- Durrant KL. Known and Hidden Sources of Caffeine in Drugs, Food, and Natural Products. *J Am Pharm Assoc* [Internet]. 2002 Jul;42(4):625–37. Available from: <http://dx.doi.org/10.1331/108658002763029607>
- Eticha S, Bedassa T. Determination of Caffeine in Coffee Samples by High-Performance Liquid Chromatography and Ultra Violet - Visible Spectrophotometry Methods from Wollega, Ethiopia. *Int J Biochem Biophys Mol Biol* [Internet]. 2020;5(1):8. Available from: <http://www.sciencepublishinggroup.com/journal/paperinfo?journalid=246&doi=10.11648/j.ijbbmb.20200501.12>
- Brunetto M del R, Gutiérrez L, Delgado Y, Gallignani M, Zambrano A, Gómez Á, et al. Determination of theobromine, theophylline, and caffeine in cocoa samples by a high-performance liquid chromatographic method with online sample cleanup in a switching-column system. *Food Chem*. 2007;100(2):459–67.
- Li S, Berger J, Hartland S. UV spectrophotometric determination of theobromine and caffeine in cocoa beans. *Anal Chim Acta*. 1990;232(C):409–12.
- Oba S, Nagata C, Nakamura K, Fujii K, Kawachi T, Takatsuka N, et al. Consumption of coffee, green tea, oolong tea,

- black tea, chocolate snacks, and caffeine content in relation to risk of diabetes in Japanese men and women. *Br J Nutr.* 2010;103(3):453–9.
24. Yalwa IR, Bello AM. Determination of caffeine content in some varieties of kola nut (*C. acuminata*). *Bayero J Pure Appl Sci.* 2018;10(1):247.
25. Vuong Q V., Roach PD. Caffeine in green tea: Its removal and isolation. *Sep Purif Rev.* 2014;43(2):155–74.
26. Vu Dang H, Truong Thi Thu H, Dong Thi Ha L, Nguyen Mai H. RP-HPLC and UV Spectrophotometric Analysis of Paracetamol, Ibuprofen, and Caffeine in Solid Pharmaceutical Dosage Forms by Derivative, Fourier, and Wavelet Transforms: A Comparison Study. *J Anal Methods Chem.* 2020;2020(c).
27. Nawrot P, Jordan S, Eastwood J, Rotstein J, Hugenholtz A, Feeley M. Effects of caffeine on human health. *Food Addit Contam.* 2003;20(1):1–30.
28. Gupta PK, Himashankari B, Sinha A. Caffeine: Benefits, Risks and Effects-A Review. *Indian J Public Heal Res Dev.* 2020;11(03):377–82.
29. Wu Q, Jiang F, Li C, Hu Y, Liu Y. Interaction of caffeine with bovine serum albumin: Determination of binding constants and the binding site by spectroscopic methods. *Chinese J Chem [Internet].* 2011 Mar;29(3):433–40. Available from: <https://onlinelibrary.wiley.com/doi/10.1002/cjoc.201190100>
30. Banipal TS, Kaur N, Banipal PK. Binding studies of caffeine and theophylline to bovine serum albumin: Calorimetric and spectroscopic approach. *J Mol Liq [Internet].* 2016;223:1048–55. Available from: <http://dx.doi.org/10.1016/j.molliq.2016.09.034>
31. Li Z, Li Z, Ma H, Fu S, Liu G, Hao C, et al. Molecular insight into binding behavior of caffeine with lactoferrin : Spectroscopic, molecular docking, and simulation study. *J Dairy Sci [Internet].* 2023;106(12):8249–61. Available from: <http://dx.doi.org/10.3168/jds.2023-23631>
32. Sovrić M, Mrkalić E, Jelić R, Serafinović MĆ, Stojanović S, Prodanović N, et al. Effect of Caffeine and Flavonoids on the Binding of Tigecycline to Human Serum Albumin: A Spectroscopic Study and Molecular Docking. *Pharmaceuticals.* 2022;15(3).
33. Topalá T, Bodoki A, Oprean L, Oprean R. Bovine serum albumin interactions with metal complexes. *Clujul Med.* 2014;87(4):5.
34. Dinu V, Borah PK, Muleya M, Scott DJ, Lithgow R, Patten J, et al. Flavor compounds affect protein structure: The effect of methyl anthranilate on bovine serum albumin conformation. *Food Chem [Internet].* 2022 Sep;388:133013.
35. Lalah JO, Wandiga SO. Extinction coefficients and dissolved organic carbon content in freshwater in Kenya. *Bull Environ Contam Toxicol.* 2006;77(4):533–42.
36. Pace CN, Vajdos F, Fee L, Grimsley G, Gray T. How to measure and predict the molar absorption coefficient of a protein. *Protein Sci.* 1995;4(11):2411–23.
37. Sudha A, Srinivasan P, Thamilarasan V, Sengottuvelan N. Exploring the binding mechanism of 5-hydroxy-3',4',7-trimethoxyflavone with bovine serum albumin: Spectroscopic and computational approach. *Spectrochim Acta A Mol Biomol Spectrosc [Internet].* 2016 Mar 15;157:170–81. Available from: <http://dx.doi.org/10.1016/j.saa.2015.12.028>
38. Ràfols C, Amézqueta S, Fuguet E, Bosch E. Molecular interactions between warfarin and human (HSA) or bovine (BSA) serum albumin evaluated by isothermal titration calorimetry (ITC), fluorescence spectrometry (FS) and frontal analysis capillary electrophoresis (FA/CE). *J Pharm Biomed Anal [Internet].* 2018;150:452–9. Available from:
- <http://dx.doi.org/10.1016/j.jpba.2017.12.008>
39. Raghav D, Mahanty S, Rathinasamy K. Characterizing the interactions of the antipsychotic drug trifluoperazine with bovine serum albumin: Probing the drug-protein and drug-drug interactions using multi-spectroscopic approaches. *Spectrochim Acta - Part A Mol Biomol Spectrosc [Internet].* 2020 Feb;226:117584. Available from: <https://linkinghub.elsevier.com/retrieve/pii/S1386142519309746>
40. Ojha H, Mishra K, Hassan MI, Chaudhury NK. Spectroscopic and isothermal titration calorimetry studies of binding interaction of ferulic acid with bovine serum albumin. *Thermochim Acta [Internet].* 2012;548:56–64. Available from: <http://dx.doi.org/10.1016/j.tca.2012.08.016>
41. Zhou KL, Pan DQ, Lou YY, Shi JH. Intermolecular interaction of fosinopril with bovine serum albumin (BSA): The multi-spectroscopic and computational investigation. *J Mol Recognit [Internet].* 2018 Aug;31(8):e2716. Available from: <https://onlinelibrary.wiley.com/doi/10.1002/jmr.2716>
42. Nienhaus K, Nienhaus GU. Probing Heme Protein-Ligand Interactions by UV/Visible Absorption Spectroscopy. *Methods Mol Biol.* 2005;305:215–41.
43. Wani TA, Bakheit AH, Abounassif MA, Zargar S. Study of interactions of an anticancer drug neratinib with bovine serum albumin: Spectroscopic and molecular docking approach. *Front Chem.* 2018;6(MAR):1–9.
44. Sen P, Fatima S, Ahmad B, Khan RH. Interactions of thioflavin T with serum albumins: Spectroscopic analyses. *Spectrochim Acta - Part A Mol Biomol Spectrosc.* 2009;74(1):94–9.
45. Lakowicz JR. Introduction to Fluorescence. In: *Principles of Fluorescence Spectroscopy.* 3rd ed. Springer; 2006. p. 1–25.
46. Wang BL, Pan DQ, Kou SB, Lin ZY, Shi JH. Exploring the binding interaction between bovine serum albumin and perindopril as well as the influence of metal ions using multi-spectroscopic, molecular docking, and DFT calculation. *Chem Phys [Internet].* 2020;530(May 2019):110641. Available from: <https://doi.org/10.1016/j.chemphys.2019.110641>
47. Maurice R. Eftink. Intrinsic Fluorescence of Proteins. In: Lakowicz JR, editor. *Topics in Fluorescence Spectroscopy [Internet].* Boston, MA: Springer US; 2000. p. 1–13. Available from: <http://link.springer.com/10.1007/b115628>
48. Wen MG, Zhang XB, Tian JN, Ni SH, Bian HD, Huang YL, et al. Binding interaction of xanthoxylin with bovine serum albumin. *J Solution Chem.* 2009;38(4):391–401.
49. Peterson FC, Anderson PJ, Berliner LJ, Brooks CL. Expression, folding, and characterization of small proteins with increasing disulfide complexity by a PT7-7-derived phagemid. *Protein Expr Purif.* 1999;15(1):16–23.
50. Belinskaia DA, Voronina PA, Batalova AA, Goncharov N V. Serum Albumin. *Encyclopedia.* 2020;1(1):65–75. 51. Raghav D, Ashraf SM, Mohan L, Rathinasamy K. Berberine Induces Toxicity in HeLa Cells through Perturbation of Microtubule Polymerization by Binding to Tubulin at a Unique Site. *Biochemistry.* 2017;56(20):2594–611.