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INFLUÊNCIA DE 1-BUTANOL, 1,2-BUTANODIOL E 1,2,3,4-BUTANOTETROL NA ADSORÇÃO DE β -LACTOGLOBULINA NA INTERFACE AR-ÁGUA

Carmen M. Romero^{1,2}, Alberto R. Albis³, Néstor E. Mendieta¹

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ABSTRACT

In this work, a systematic study on the effect of 1-butanol, 1,2-butanediol and 1,2,3,4-butanetetrol (erythritol) on the surface tension of β -lactoglobulin in aqueous solution at pH 6.5 and 298.15 K is presented.

The experimental data were used to calculate the surface pressure and were adjusted to different protein adsorption models at the liquid-air interface to explain the behavior of β -lactoglobulin in aqueous solution. The results show that the alcohols produce a significant effect on the adsorption behavior of the protein at the interface that is related to the number of hydroxyl groups.

Key words: β -lactoglobulin, 1-butanol; 1,2-butanediol; 1,2,3,4-butanetetrol, surface tension, adsorption.

RESUMEN

En este trabajo se presenta un estudio sistemático del efecto de 1-butanol, 1,2-butanediol y 1,2,3,4-butanotetrol (eritritol) sobre la tensión superficial de la β -lactoglobulina en solución acuosa a 298,15 K.

Los datos experimentales fueron usados para calcular la presión superficial y se ajustaron a distintos modelos de adsorción en la interfase líquido-aire para explicar el comportamiento de la β -lactoglobulina en solución acuosa. Los resultados muestran

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que los alcoholes tienen un efecto significativo en el proceso de adsorción de la proteína en la interfase, relacionado con el número de grupos hidroxilo del alcohol.

Palabras clave: β -lactoglobulina, 1-butanol; 1,2-butanodiol; 1,2,3,4-butanotetrol; tensión superficial, adsorción.

RESUMO

Neste trabalho, um estudo sistemático sobre o efeito do 1-butanol, 1,2-butanodiol e 1,2,3,4-butanotetrol (eritritol) sobre a tensão superficial da β -lactoglobulina em solução aquosa em pH 6,5 e 298,15 K é apresentado. Os dados experimentais foram utilizados para calcular a pressão de superfície e foram ajustados para diferentes modelos de adsorção de proteínas na interface líquido-ar para explicar o comportamento de β -lactoglobulina em solução aquosa. Os resultados mostram que os alcoóis produzem um efeito significativo sobre o comportamento de adsorção da proteína na interface que está relacionado com o número de grupos hidroxila.

Palavras-chave: β -lactoglobulina, 1-butanol; 1,2-butanodiol; 1,2,3,4-butanotetrol; tensão superficial; adsorção.

INTRODUCTION

Protein stability is a consequence of a delicate balance between intermolecular interactions of the protein with solvent molecules, which are dominant at high dilution and intramolecular interactions between the functional groups of the protein. While most researchers agree that the hydrophobic effect plays a key role in stabilizing proteins, there is not a defi-

nite explanation concerning whether and to what extent a given type of interaction determines the native conformation of a protein (1-3).

Several proposals have been presented to explain the effect of cosolvents on protein stability. Between them, some authors propose that the stabilizing effect of osmolytes such as polyols and sugars is a consequence of the increase in the surface tension of the solvent (1-3), based on experimental results that suggest that, with few exceptions, additives that increase the surface tension of water also stabilize proteins (4). However other studies do not show a linear dependence of denaturation temperatures with the increase in the solvent surface tension (5,6) and the correlation between thermal stabilization and the change in surface tension of the solvent remains to be an unresolved problem that requires the knowledge of the adsorption characteristics of proteins and is worth of systematic research using well characterized model macromolecules.

β -Lactoglobulin has been considered a model globular protein (7-13). It is one of the most investigated proteins and its structural and dynamic properties have been extensively studied. β -Lactoglobulin is one of the main whey proteins and has several applications in food and pharmaceutical industries. In its native state is a globular protein (7-10) with a molecular weight of 18362 g mol⁻¹, 162 amino acid residues, two disulfide bonds, and an isoelectric point around 5.2. X Ray diffraction and NMR studies show that it is a predominantly β -sheet protein; the secondary structure of this protein consists of 9 strands of beta structure,

an α -helix segment and three helicoidal turns (11-14). It has been reported that between pH 5.2 and 7.5 β -lactoglobulin exists in a dimeric form while under pH 3.0 and above pH 8.0 the protein exists as a monomer (15). However, other results clearly show that near neutral pH the monomeric form of β -lactoglobulin is dominant (16).

Several studies on the influence of alcohols and polyols on β -lactoglobulin thermal stability have been developed. The results show that polyols improve the conformational stability of proteins while alcohols induce protein denaturation and the effect decreases as the number of OH groups increases (16-18).

Proteins are surface active substances that lower surface tension of water and tend to adsorb at the water-air interface. Adsorption of β -lactoglobulin in aqueous solution has been studied in buffers (17,19,20) and in the presence of different cosolvents such as denaturing agents (20), polysaccharides (21), and polyols (22). The effect of sugars has been explained as a consequence in the increase in the surface tension of water considering it the major factor in the stabilization of proteins. In the case of polyols, some authors suggest that surface tension of water does not have a major effect in protein stabilization and other effects such as preferential hydration and solvophobic effects are considered to be the responsible for protein stabilization (23-24) while other studies suggest that the surface tension of water has a fundamental role (25).

Protein adsorption is a complex dynamic process that is affected by protein

structure, intermolecular forces between the adsorbed molecules and the solvent, solute-solute and solvent-solvent interactions and the presence of other substances (25-29). Dynamic surface tension is very sensitive to adsorption. As the protein is adsorbed at the liquid-air interface, the surface tension decreases until the equilibrium value is attained. Thus the process can be followed from time evolution of surface pressure π defined as the difference between the surface tension of the protein solution γ and the surface tension of the solvent γ^0 :

$$\pi = \gamma - \gamma^0 \quad [1]$$

Adsorption involves different processes before equilibrium surface tension is attained. Two empirical models have been proposed to describe protein adsorption. One of the models involves two kinetic surface tension regimes and the other three regimes, being the first step in both of them the diffusion of the protein from the bulk toward the interface. Additionally, in some cases an induction period is observed in which the surface tension remains nearly equal to that of the pure solvent.

The diffusion controlled step is usually represented by equation (21, 30-33):

$$\pi = 2k C_0 T \left(\frac{Dt}{3.14} \right)^{1/2} = A t^{1/2} \quad [2]$$

where k is the Boltzmann constant, C_0 is the protein concentration in the bulk, D is the diffusion coefficient of the protein in the solvent, T is the absolute temperature, A is a constant and t is the drop lifetime at which the surface pressure π is measured.

After some time, protein concentration at the interface increases and the rate of the process is controlled by the adsorption and rearrangement of adsorbed molecules. At this stage, the exposure of some part of the protein to the air can produce reorientation and conformational changes in the protein. This process can be represented by one or two steps that are usually represented by semi empirical first-order equations (30, 34).

$$\pi = \pi_f - \pi_0 e^{-k(t-t_0)} \quad [3]$$

where π_f , π_0 and π are the surface pressures at the final adsorption time of each step, at the initial time t_0 and at any time t of drop formation, and k is the first-order kinetic constant.

For small globular proteins at low concentration, adsorption at the air-water interface usually follows a two steps model: Diffusion and rearrangement. However, the adsorption rate depends also on the nature of the solvent (pH, cosolvents) and factors such as temperature and pressure, so the adsorption model has to be determined from experimental behavior of the protein.

In the present work, the influence of the cosolvents 1-butanol, 1,2-butanediol and 1,2,3,4-butanetriol (erythritol) on the dynamic surface tension of β -lactoglobulin in aqueous solution at 298.15 K and pH 6.5 is considered. Time evolution of surface pressure is used to analyze the effect of the number of hydroxyl groups of the alcohols on the adsorption dynamics of the protein.

MATERIALS AND METHODS

The materials used in this work were the following: β -lactoglobulin 90 % (Sigma), 1-butanol ≥ 99.5 % (Merck), 1,2-butanediol 98 % (Aldrich), and 1,2,3,4-butanetriol (erythritol) ≥ 99 % (Sigma). Water content of the alcohols and liquid polyols was determined by the Karl Fischer's method, and they were degassed before used. Water was doubly distilled, treated according to literature (34), and degassed before used to obtain water with conductivity lower than $2 \mu\text{S}\cdot\text{cm}^{-1}$.

Solutions were prepared by weight using a Mettler balance AT 201 with sensitivity of 10^{-5} g in the lower range. Polyol solutions were prepared in molar fractions x_{OH} of 0.005; 0.010; 0.015, and 0.020 but for butanol the highest molar fraction used was 0.015 due to its low solubility. Solutions of β -lactoglobulin (BLG) with mass fractions w_{BLG} of $9.86 \cdot 10^{-5}$ and $6.39 \cdot 10^{-5}$ were prepared dissolving the protein in the aqueous solutions. The final pH in all cases was 6.5.

Surface tension γ measurements were determined using a LAUDA TVT2 drop volume tensiometer based on the principle of the pending drop volume, with temperature control better than 0.01 K and uncertainty of $\pm 0.1 \text{ mN}\cdot\text{m}^{-1}$ in surface tension. The volume of the syringe used for the measurements was 2.5 mL and the tip used has an external diameter of 1.395 mm. Surface tension of a solution of β -lactoglobulin in water of mass fraction $9.86 \cdot 10^{-5}$ and the aqueous polyol solutions were measured using the dynamic method with a time of drop for-

mation of 60 s and reported values are the average of 18 to 20 measurements. The surface tension of β -lactoglobulin with mass fractions of $9.86 \cdot 10^{-5}$ and $6.39 \cdot 10^{-5}$ in the aqueous solvents was measured using the quasi-static method (35-37). The surface tension value corresponds to the average of at least three independent measurements.

Density of solutions was measured using an Anton Paar vibrating tube densimeter DSA 5000 calibrated using dry air and distilled water at $298.15\text{K} \pm 0.01\text{K}$. The uncertainty in density measurements is $\pm 5 \cdot 10^{-6} \text{g} \cdot \text{cm}^{-3}$.

RESULTS AND DISCUSSION

Experimental data obtained in this work for equilibrium surface tension γ at 298.15K of aqueous solutions of 1-butanol, 1,2-butanediol and 1,2,3,4-butanetetrol as a function of mole fraction x_{OH} , and the values for aqueous solutions of polyols in the presence of β -lactoglobulin at $9.84 \cdot 10^{-5}$ mass fraction w_{BLG} are presented in Table 1.

The results obtained for surface tension of butanol and 1,2,3,4-butanetetrol in water agree well with literature data (36, 38, 39) but the value reported for 1,2-butanediol is lower than the results obtained in this work (39).

From the experimental data it can be observed that 1-butanol and 1,2-butanediol lead to a decrease in the surface tension of water being the larger change observed with butanol. The addition of 1,2,3,4-butanetetrol induces a very small change in surface tension and shows a

complex behavior that does not follow a clear trend.

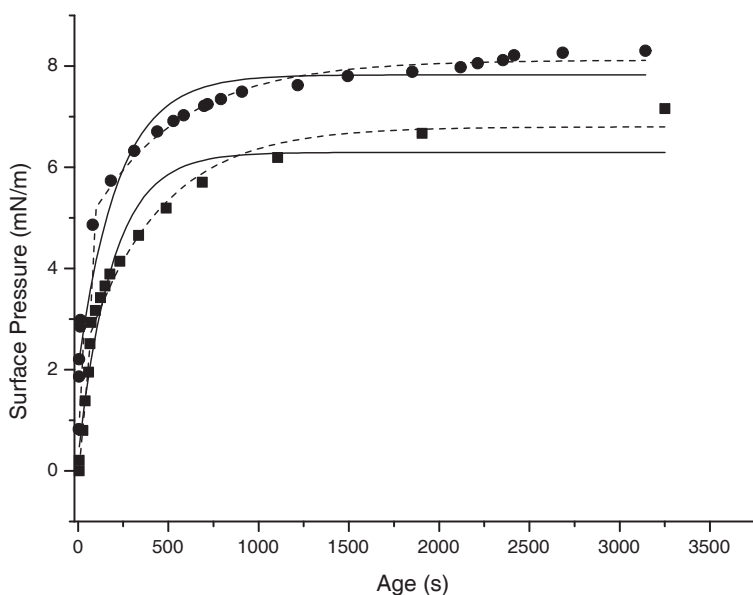
The results in Table 1 show the effect of β -lactoglobulin at $9.86 \cdot 10^{-5}$ mass fraction on the equilibrium surface tension of the aqueous solutions of polyols. As expected, the protein lowers the surface tension of water indicating that protein molecules tend to adsorb at the interface due to its amphiphilic nature giving as result a positive surface excess. The decrease in the surface tension is larger in the presence of 1-butanol followed by 1,2-butanediol while 1,2,3,4-butanetetrol produces a small change in surface tension and shows a complex behavior that does not follow a clear trend. A similar behavior has been reported for β -lactoglobulin in the presence of sorbitol (22).

Protein adsorption was followed measuring the change of surface tension as a function of time using two different mass fractions of protein: $9.86 \cdot 10^{-5}$ and $6.39 \cdot 10^{-5}$, respectively. The experimental protein adsorption profiles were adjusted to the two and three steps models using the Origin® software.

The results are shown in Figure 1 for the selected protein concentrations. Surface pressure increases as protein concentration becomes larger and induction period is not observed before the adsorption process. The time dependence of the change in surface pressure depends on protein concentration and using dynamic surface measurements it has been shown that for low concentrations of β -lactoglobulin an induction period is observed while it is not present at higher protein concentrations (40, 41).

Table 1. Equilibrium surface tension of butanol, 1,2-butanediol, and 1,2,3,4-butanetriol in water and aqueous solutions of β -lactoglobulin at 298.15 K and pH 6.5.

Polyol-water		Polyol-water- β -lactoglobulin		
X_{OH}	$\gamma/mN \cdot m^{-1}$	X_{OH}	w_{BLG}	$\gamma/mN \cdot m^{-1}$
0.000	72.00	0.000	$9.86 \cdot 10^{-5}$	63.68
1-butanol				
0.005	44.10	0.005	$9.86 \cdot 10^{-5}$	35.06
0.010	35.81	0.010	$9.86 \cdot 10^{-5}$	27.16
0.015	29.15	0.015	$9.86 \cdot 10^{-5}$	20.64
1,2-butanediol				
0.005	59.55	0.005	$9.86 \cdot 10^{-5}$	50.77
0.010	56.13	0.010	$9.86 \cdot 10^{-5}$	47.34
0.015	53.89	0.015	$9.86 \cdot 10^{-5}$	44.75
0.020	50.28	0.020	$9.86 \cdot 10^{-5}$	40.45
1,2,3,4-butanetriol				
0.005	71.16	0.005	$9.86 \cdot 10^{-5}$	58.49
0.010	71.91	0.010	$9.86 \cdot 10^{-5}$	57.84
0.015	72.48	0.015	$9.86 \cdot 10^{-5}$	57.98
0.020	73.13	0.020	$9.86 \cdot 10^{-5}$	58.56


Figure 1. Surface pressure β -lactoglobulin in water at 298.15 K. Protein mass fraction: 6.39×10^{-5} (■), 9.86×10^{-5} (●). Two step model (—), three step model (- - -)

Experimental data were fitted to the two general models and they are well described by a three step model. After a rapid diffusion step the rate of β -lactoglobulin adsorption is controlled by the penetration and by rearrangement and partial unfolding of the protein at the interface. The diffusion controlled step is too rapid to be measured accurately using the quasi-static method (less than 30 s) and the other two steps follow a logarithmic behavior.

Table 2 shows the adsorption parameters of the two logarithmic steps of the model. π_1 and k_1 are the final pressure and the kinetic constant of the penetration process and π_2 and k_2 are the final pressure and the kinetic constant associated to the protein rearrangement at the liquid-air interface.

The results presented reveal that the constants for the penetration and for the rearrangement steps are nearly the same for the two protein concentrations and that rate of the penetration process is much higher than the rate of rearrangement of β -lactoglobulin at the liquid-air interface.

Adsorption at the air-water interface in the presence of alcohols was determined following the change of surface tension as a function of time for solutions of β -lactoglobulin of mass fraction of $9.86 \cdot 10^{-5}$ in the aqueous solvents at 298.15 K and pH 6.5. The effect of al-

cohol concentration on the adsorption behavior of the protein is shown in Figure 2. The presence of alcohols increases surface pressure when compared with the behavior of the protein in water and the effect is larger as alcohol concentration increases. The change in surface pressure for polyols depends clearly on the number of OH groups. The largest increase is observed with 1,2,3,4-butanetriol followed by 1,2-butanediol and the smallest change is observed in the presence of 1-butanol.

The experimental protein adsorption profile in the presence of 1-butanol, 1,2-butanediol and 1,2,3,4-butanetriol was adjusted to the two and three steps models. In the presence of alcohols, the experimental data fitted well the two steps model, as it is shown in Figure 2 indicating that the three alcohols induce an important modification in the adsorption behavior of the protein. The diffusion controlled step is fast and occurs below the lowest experimental time measured as for the protein in water. After the diffusion stage, the observed behavior shows that adsorption of β -lactoglobulin is controlled by penetration of β -lactoglobulin at the liquid-air interface. The absence of the rearrangement regime suggests that the conformational changes in protein structure are very rapid and depend on the nature of the cosolvent. Table 3 shows the adsorption parameters of the two steps model.

Table 2. Value of the adsorption parameters for β -lactoglobulin in water at 298.15K

w_{BLG}	π_1 (mN·m ⁻¹)	π_2 (mN·m ⁻¹)	k_1 (s ⁻¹)	k_2 (s ⁻¹)
$6,39 \cdot 10^{-5}$	4.4	6.7	$9.1 \cdot 10^{-3}$	$2.7 \cdot 10^{-3}$
$9,86 \cdot 10^{-5}$	4.6	8.1	$13.8 \cdot 10^{-3}$	$2.5 \cdot 10^{-3}$

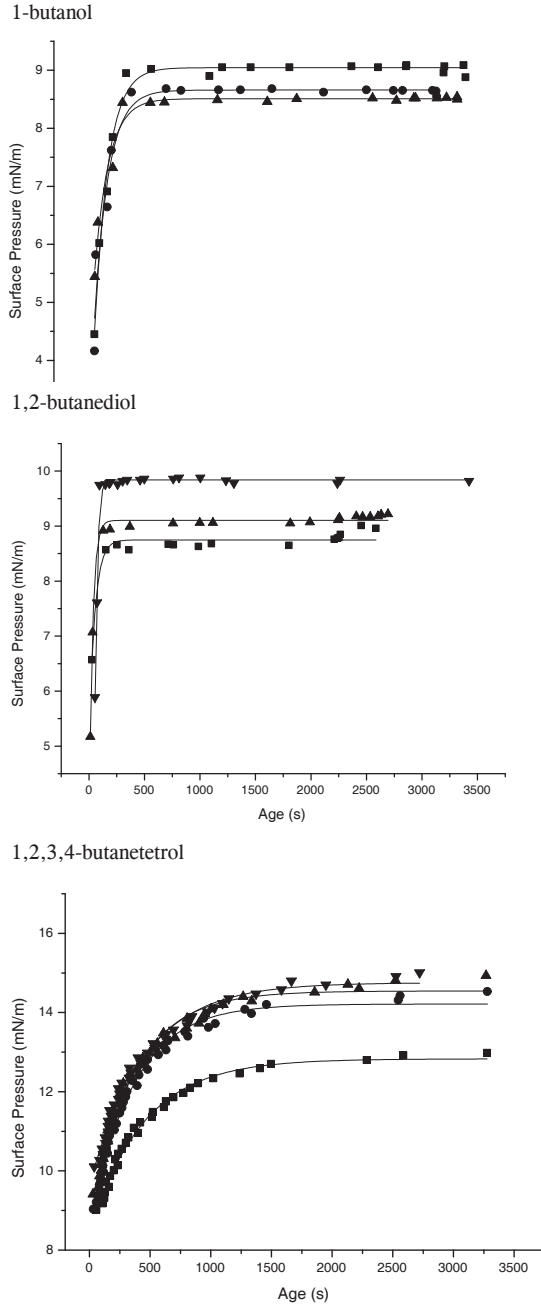


Figure 2. Effect of alcohol concentration on the adsorption profiles of β -lactoglobulin at pH 6.5 and 298.15 K. Alcohol mole fraction: 0.0050 (■), 0.010 (●), 0.015 (▲), 0.020 (▼).

Table 3. Value of the adsorption parameters for β -lactoglobulin in the presence of alcohols at 298.15 K.

Cosolvent mole fraction	Adsorption step		Rearrangement step	
	π_1 mN·m ⁻¹	k_1 s ⁻¹	π_2 mN·m ⁻¹	k_2 s ⁻¹
1-butanol				
0.0000	4.6	1.3×10^{-3}	8.1	2.5×10^{-3}
0.0050	9.0	1.2×10^{-2}		
0.0100	8.7	1.1×10^{-2}		
0.0150	8.5	1.1×10^{-2}		
1,2-butanediol				
0.0000	4.6	1.3×10^{-3}	8.1	2.5×10^{-3}
0.0050	8.7	1.9×10^{-2}		
0.0100	8.8	1.6×10^{-2}		
0.0150	9.1	3.2×10^{-2}		
0.0200	9.8	4.6×10^{-2}		
1,2,3,4-butanetetrol				
0.0000	4.6	1.3×10^{-3}	8.1	2.5×10^{-3}
0.0050	12.8	2.3×10^{-3}		
0.0100	14.2	2.8×10^{-3}		
0.0150	14.5	2.7×10^{-3}		
0.0200	14.8	2.3×10^{-3}		

The equilibrium surface pressure π_1 for the penetration step increases with the addition of butanol and butanediol being the largest change observed in the presence of 1,2,3,4-butanetetrol. This behavior shows that the increase in surface pressure is largest for the more hydrophilic and non surface active cosolvent, while for the hydrophobic and surface active compounds butanol and butanediol, the change is smaller. In the case of butanol, after an initial increase, the surface pressure tends to become lower as the concentration of the cosolvent increases.

The kinetic constants follow a different behavior with cosolvent concentration. The presence of butanol induces a fast initial increase of k_1 and at higher concentration its value remains constant and the kinetic constant for butanediol becomes larger as concentration increases. This behavior suggests that protein conformation changes are faster in the presence of butanol, which has a strongest denaturing effect. 1,2,3,4-butanetetrol causes a very small change of the constant indicating that the adsorption regime of the protein is not affected by the pol-

yol because it is preferentially excluded from the liquid-air interface and does not induce changes in protein conformation.

CONCLUSIONS

In the present work, a systematic study on the effect of 1-butanol, 1,2-butanediol and 1,2,3,4-butanetetrol on the surface tension of β -lactoglobulin in aqueous solution at pH 6.5 and 298.15 K is presented. The protein lowers the surface tension of water indicating that protein molecules adsorb at the interface giving as result a positive surface excess. The decrease in the equilibrium surface tension is larger in the presence of 1-butanol followed by 1,2-butanediol while 1,2,3,4-butanetetrol produces a small change.

The adsorption profile of β -lactoglobulin in water, determined from the dynamic surface pressure, is well described by a three step model. An initial and rapid diffusion step followed by the penetration regime and the rearrangement and partial unfolding step of the protein at the interface. The presence of the alcohols induces important changes in the adsorption behavior of the protein. The diffusion controlled step is fast as for the protein in water and after the diffusion stage, adsorption of β -lactoglobulin is controlled by penetration of β -lactoglobulin at the liquid-air interface. The absence of the rearrangement regime suggests that if the alcohols induce conformational modifications in protein structure, they are very rapid. The adsorption behavior of the protein suggests that protein conformation changes are faster in the presence of butanol, while 1,2,3,4-butanetetrol does not induce changes in protein native structure.

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