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iournal or	lournal of bioscience and bioengineering
publication title	
volume	11/
number	5
page range	539-543
year	2014-05
権利	(C) 2013, The Society for Biotechnology,
	Japan. NOTICE: this is the author's version
	of a work that was accepted for publication in
	Journal of bioscience and bioengineering
	Changes resulting from the publishing process
	changes resulting from the publishing process,
	such as peer review, editing, corrections,
	structural formatting, and other quality
	control mechanisms may not be reflected in
	this document. Changes may have been made to
	this work since it was submitted for
	publication. A definitive version was
	subsequently published in Journal of
	bioscience and bioengineering, 117, 5, 2014
	DOI: 10.1016/j.jbiosc.2013.10.016
URL	http://hdl.handle.net/2241/00121920

doi: 10.1016/j.jbiosc.2013.10.016

Arginine and Lysine Reduce the High Viscosity of
Serum Albumin Solutions for Pharmaceutical Injection
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Running title: Reducing viscosity with Arg and Lys

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19 Abstract

20 Therapeutic protein solutions for subcutaneous injection must be very highly concentrated, which 21 increases their viscosity through protein-protein interactions. However, maintaining a solution 22 viscosity below 50 cP is important for the preparation and injection of therapeutic protein 23 solutions. In this study, we examined the effect of various amino acids on the solution viscosity 24 of very highly concentrated bovine serum albumin (BSA) and human serum albumin (HSA) at a 25 physiological pH. Among the amino acids tested, L-arginine hydrochloride (ArgHCl) and L-26 lysine hydrochloride (LysHCl) (50-200 mM) successfully reduced the viscosity of both BSA and 27 HSA solutions; guanidine hydrochloride (GdnHCl), NaCl, and other sodium salts were equally as 28 effective, indicating the electrostatic shielding effect of these additives. Fourier transform 29 infrared spectroscopy showed that BSA is in its native state even in the presence of ArgHCl, 30 LysHCl, and NaCl at high protein concentrations. These results indicate that weakened protein-31 protein interactions play a key role in reducing solution viscosity. ArgHCl and LysHCl, which 32 are also non-toxic compounds, will be used as additives to reduce the solution viscosity of 33 concentrated therapeutic proteins.

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35 Keywords: viscosity, arginine, lysine, serum albumin, protein-protein interaction

37 Introduction

38 Therapeutic proteins account for an increasingly large proportion of pharmaceutical drugs 39 in recent years. These proteins are normally administered by subcutaneous injection (1). When a 40 high dose of therapeutic proteins is required, the protein solution must often be very highly 41 concentrated to reduce the injection volume for patient convenience. However, such a high 42 protein concentration frequently leads to high viscosity, posing considerable challenges for both 43 processing and injection (2-4). For example, subcutaneous injection is generally performed with 44 solutions under 50 cP to reduce the time required for injection (5). Therefore, maintaining a 45 solution viscosity below 50 cP is a primary goal when developing high concentration 46 formulations of therapeutic proteins.

47 The viscosity of a protein solution is caused by noncovalent protein-protein interactions 48 such as electrostatic attraction or repulsion, leading to the formation of a transient three-49 dimensional network of protein (6). There has been considerable effort directed toward lowering 50 the viscosity of concentrated protein solutions through the application of various solution 51 additives (7-11). Among them, inorganic salts (12, 13) and hydrophobic salts (5, 14) have been 52 shown to effectively reduce the viscosity of protein solutions. However, some of these salts have 53 adverse effects on proteins, leading to destabilization and consequent aggregate formation (15). 54 Thus, there is always a demand for effective formulation conditions that reduce the viscosity of a 55 solution without compromising protein stability.

We have developed the application of ArgHCl for suppressing aggregation and adsorption of various proteins during heat treatment and oxidative refolding (16–22). These effects have been ascribed, at least in part, to the interaction of aromatic groups on arginine with the protein surface (23). We have shown that ArgHCl binds to aromatic amino acids in proteins

60 using a crystal structure (24) and increases the solubility of low molecular weight solutes 61 containing aromatic moieties, consistent with the above mechanism (25–29). ArgHCl does not 62 destabilize the protein structure but decreases the probability of aggregation (30–36). Based on 63 its effects on protein aggregation, ArgHCl might also reduce the viscosity of protein solutions by 64 suppressing transient protein-protein interactions, thus preventing the formation of a threedimensional network. In this work, we examined the effect of various amino acids on the 65 66 viscosity of bovine and human serum albumin (BSA and HSA, respectively). HSA is a clinically 67 important supplement for the loss of body fluid. This is the first report describing the application 68 of amino acids for reducing the viscosity of HSA solutions.

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71 Material and Methods

BSA and HSA were obtained from Sigma Chemical Co. (St. Louis, MO). The purity of BSA and HSA was more than 98% and 96%, respectively. L-Alanine (Ala), L-valine (Val), Lmethionine (Met), L-proline (Pro), glycine (Gly), L-serine (Ser), L-threonine (Thr), L-lysine hydrochloride (LysHCl), L-arginine hydrochloride (ArgHCl), L-histidine (His), guanidine hydrochloride (GdnHCl), sodium chloride (NaCl) were obtained from Wako Pure Chemical Inc. Ltd. (Osaka, Japan). All chemicals were of reagent grade and were used as received.

The sample solution was prepared as follows: Serum albumin was dissolved in deionized water, and the pH of the protein solution was adjusted to the desired values with 0.5 M NaOH or HCl. The additive solution was mixed with the serum albumin solution to achieve the appropriate concentrations of protein and additive, and the pH was then readjusted. The viscosity of the sample solution was measured with a torsional oscillation VM-10A-L viscometer (CBC Materials Co. Ltd., Japan) at 25°C without depending on shear rate. A 1 ml sample was loaded onto the measuring microtube. The viscosity of each sample was measured at least in duplicate. Error bars in Figures depict the standard deviation of the mean of three independent experiments.

Fourier transform infrared (FTIR) spectra of the sample solutions were measured on a FT-IR 4200 spectrometer (JASCO Corporation, Japan) using the attenuated total reflection (ATR) method. The sample solution was placed on the surface of a single-reflection ZnSe prism on an ATR accessory. Spectra were collected in the range of 1,700 to 1,500 cm⁻¹ with a spectral resolution of 2.0 cm⁻¹.

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94 **Results and Discussion**

95 **Examination of the solution conditions**

96 The viscosity of a protein solution is due, at least in part, to protein-protein interactions 97 and is therefore expected to change with pH; this behavior is similar to protein solubility, which 98 also depends on solution pH. The viscosity of a 40 mg/ml BSA solution as a function of pH was reported previously (37). The lowest viscosity was observed at around pH 4.8, near the isoelectric 99 100 point (pI) of BSA, which is in good agreement with our experimental value (data not shown). It 101 is clear that the viscosity is greater when BSA is positively charged at pH 4.0 and negatively 102 charged above pH 6.0, which suggests the involvement of charge-charge repulsive interactions in 103 network formation. Figure 1 shows that the viscosity of the BSA solution at pH 7.4 is a function of the protein concentration up to 300 mg/ml. The solution viscosity remained nearly constant
below 150 mg/ml. Further increases in protein concentration resulted in an exponential increase
in viscosity to over 100 cP at 300 mg/ml.

107 In order to confirm the structural change by such high concentration, we measured FTIR 108 spectra of BSA as a function of protein concentration at pH 6.8, which is the pH value of BSA 109 solutions not adjusted, due to the avoidance of unnecessary factors (Fig. 2). The amide I and 110 amide II regions from 1,700 to 1,500 cm⁻¹ of the FTIR spectra were clearly observed with the 111 BSA concentration of 75-300 mg/ml, which reflects the second structure of protein. The maximal 112 intensity of the amide I region at 1651 cm⁻¹ increased with increasing the protein concentration. 113 The normalized spectra of the amide I and amide II regions of BSA with the concentration of 75-114 300 mg/ml were identical (Fig. 2B), indicating that the secondary structure of BSA retained 115 constant even in the presence of high protein concentration.

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117 Effect of amino acids on the viscosity of BSA solutions

118 Because the viscosity of BSA solution was dependent on both pH and protein 119 concentration, we took these parameters into consideration when determining the solution 120 conditions for evaluating the effects of various additives. We chose a pH of 7.4 because it is 121 preferable for injectable solutions to be similar to physiological conditions. At this pH, a 275 122 mg/ml BSA solution showed a viscosity of ~50 cP (see Fig. 1). Any additives that lower the 123 viscosity below this value may be acceptable for pharmaceutical applications (e.g., subcutaneous 124 injection). Using a pH of 7.4 and a concentration of 275 mg/ml, we examined the effects of 125 additives on the viscosity of a BSA solution.

126 Figure 3 shows the viscosity of a 275 mg/ml BSA solution at pH 7.4 in the presence of 127 200 mM additives. The additives tested were the 10 naturally occurring amino acids shown in 128 Fig. 3; other amino acids were insufficiently soluble in the presence of 275 mg/ml BSA. Other 129 additives were also tested for control experiment; NaCl served as salts, and GdnHCl served as 130 ArgHCl. Errors in the viscosity measurements may be mostly attributed to preparation errors 131 such as alterations in protein concentration because no viscosity errors were noted when the same 132 sample solution was measured. As shown in Fig. 3, the viscosity of the BSA solution was ~50 cP 133 in the absence of additives. In the presence of 200 mM ArgHCl or LysHCl, the viscosity of the 134 BSA solution decreased to 28 cP and 25 cP, respectively, corresponding to 1.9-fold and 2.0-fold 135 respective reductions in viscosity relative to the solution without additives. Conversely, the other 136 amino acids barely reduced the viscosities of the BSA solutions. The effects of NaCl and 137 GdnHCl on viscosity were the same as those of ArgHCl and LysHCl. The observed similarity 138 between LysHCl, ArgHCl, NaCl, and GdnHCl suggests that the ionic properties of these 139 electrolytes may play an important role in reducing the viscosity of BSA solutions, as all of these 140 electrolytes are monovalent ions at neutral pH.

Figure 4 shows the viscosity of 275 mg/ml BSA in the presence of 0 to 200 mM LysHCl, ArgHCl, and NaCl. The viscosity of BSA sharply decreased with increasing concentrations of ArgHCl, LysHCl, and NaCl up to ~50 mM. Above 50 mM, the viscosity of BSA appeared to reach a plateau and was nearly independent of the additives, with all additives leading to a 1.5fold reduction in viscosity. FTIR spectra showed that the secondary structure of BSA was not altered even in the presence of 200 mM ArgHCl, LysHCl, and NaCl (Fig. 5). These data indicate that BSA does not undergo structural changes during reduction of the solution viscosity.

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149 Effect of amino acids on the viscosity of HSA solutions

150 Finally, we performed the same experiments with human serum albumin (HSA) which is 151 similar in the pI and the sequence to BSA, as it has commercial value as a formulation excipient 152 and plasma expander. Figure 6 shows the viscosity of 305 mg/ml HSA at pH 7.4 in the presence 153 of 200 mM amino acid or salt. A pattern similar to that observed for BSA was noted; the 154 viscosity of HSA was decreased 1.6-fold by ArgHCl and LysHCl, whereas the other amino acids 155 had marginal effects. As with BSA, NaCl and GdnHCl were as effective as, or slightly more 156 effective than, LysHCl and ArgHCl with respect to reducing the viscosity of the HSA solution.

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Molecular mechanisms of viscosity reduction by ArgHCl and LysHCl 158

159 The pH dependence of BSA solution viscosity is intriguing. The viscosity was at a 160 minimum at approximately pH 4.8, which is near the pI of BSA (37). This result indicated that 161 the protein-protein interactions in relation to solution viscosity were also at a minimum. In other 162 words, the viscosity of BSA solutions significantly increased on both sides of the pI, meaning 163 that stronger protein-protein interactions that affect solution viscosity occur when BSA acquires 164 net charges, regardless of whether these charges are negative or positive. Although the 165 acquisition of net charges causes overall repulsion between protein molecules, either 166 deprotonation of carboxyl groups (below the pI) or protonation of His and Lys (above the pI) 167 may enhance the local electrostatic interaction between BSA molecules, leading to increased 168 formation of transient noncovalent networks and a consequent increase in solution viscosity.

169 We expected that protein solubility would be related to solution viscosity. Protein 170 solubility is normally minimal at the pI (38). Near the pI, the net charge of a protein should be 171 close to zero, causing minimal charge repulsion and enhancing protein-protein interactions. 172 Protein-protein interactions that determine protein solubility may be related to the transient formation of a protein network that increases solution viscosity. Thus, it was expected that the viscosity of BSA solution would be maximal near its p*I*. However, the viscosity was at a minimum near the p*I* of BSA (37). Electrostatic interactions are thought to be important in high concentration formulations of monoclonal antibodies (13, 39). Thus, this unexpected result indicated that the type of protein-protein interaction that plays a role in high viscosity is different from the type of interaction responsible for determining protein solubility.

179 The effects of additives were examined at the physiological pH of 7.4. At this pH, BSA 180 has a net negative charge that apparently results in a greater viscosity than that which occurs at 181 the isoelectric pH of ~5 (37). The current study revealed that ArgHCl or LysHCl at 200 mM 182 significantly reduced the viscosities of both BSA and HSA solutions, although those of 50 mM 183 may be sufficient to achieve the same level of reduction (Fig. 4). The viscosity of albumin 184 solution was 1.6-fold lower in the presence of ArgHCl or LysHCl than in their absence. A 1.6-185 fold reduction in viscosity (from 50 cP to 30 cP) triggered by 200 mM ArgHCl or LysHCl at a 186 physiological pH may be sufficient for injection. To our knowledge, this is the first report on the 187 use of amino acids to reduce the viscosity of albumin solutions under practical conditions.

188 The molecular mechanisms by which ArgHCl and LysHCl reduce solution viscosity will 189 be discussed in detail below. It is evident that LysHCl and ArgHCl as well as their salts are 190 effective for reducing the viscosity of BSA and HSA solutions. In other words, there is no 191 specific property of Lys or Arg that plays a key role in BSA and HSA solution viscosity. We 192 have suggested the importance of electrostatic interactions in determining the protein-protein 193 network, which is closely related to the high viscosity of BSA and HSA solutions. It has been 194 reported that salts can effectively weaken such electrostatic interactions (40). Thus, both LysHCl 195 and ArgHCl may be working simply as salts. This may be the first rare case in which ArgHCl is 196 not more effective at suppressing protein-protein interactions compared with LysHCl or salts,

197 which in turn suggests less involvement of aromatic or hydrophobic interactions in the viscosity 198 of BSA and HSA solutions. When these types of interactions are involved (e.g., protein-protein 199 interactions that determine solubility or aggregation), ArgHCl normally exerts stronger effects 200 than any of the other additives tested here.

201 In conclusion, we examined the effect of amino acids on the viscosity of highly 202 concentrated solutions of BSA and HSA. We showed that (i) ArgHCl and LysHCl are effective at 203 reducing the viscosity of both BSA and HSA solutions, (ii) GdnHCl and NaCl equally reduce the 204 solution viscosity of BSA, and (iii) the ionic properties of these additives play a key role in 205 reducing the viscosity of protein solutions. These results suggested that both ArgHCl and LysHCl 206 maybe working simply as salts, and can effectively weaken electrostatic interaction. It should be 207 noted that ArgHCl is one of the most used solution additive that decreases the probability of 208 protein aggregation (30-36), adsorption on solid surface (16), and increase the solubility of 209 water-insoluble drugs (27-29) and reduced-denatured proteins (19, 20). This is the first report 210 that amino acids reduce the solution viscosity of HSA, which is a clinically important product 211 that is used as a supplement for the loss of body fluid.

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214 Acknowledgments

This work was supported by a Grant-in-Aid from JSPS KAKENHI (Grant Number 216 23550189) and by the JSPS Research Fellowships from the Ministry of Education, Culture, 217 Sports, Science, and Technology (MEXT), Japan.

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329 Figure legends

Figure 1. Viscosity of BSA solutions as a function of the protein concentration at pH 7.4.
Various concentrations of BSA were prepared at pH 7.4 and 25°C.

332

Figure 2. FTIR spectra in the amide I and amide II regions of BSA solutions as a function of the protein concentration. Original (A) and normalized (B) spectra of BSA at pH 6.8 and 25°C: 75 mg/ml (dotted line), 150 mg/ml (broken and dotted line), 225 mg/ml (broken line) and 300 mg/ml (solid line).

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Figure 3. Viscosity of BSA solutions in the absence or presence of additives. Samples containing
275 mg/ml BSA with 200 mM amino acid or salt were prepared at pH 7.4 and 25°C. Error bars
depict the standard deviation of the mean of three independent experiments.

341

Figure 4. Viscosity of BSA solutions as a function of additive concentration. Samples containing
275 mg/ml BSA with various concentrations of LysHCl (closed circle), ArgHCl (open square)
and NaCl (closed triangle) were prepared at pH 7.4 and 25°C.

Figure 5. FTIR spectra in the amide I and amide II regions of BSA solutions with additives.
Samples containing 275 mg/ml BSA with 200 mM additives were prepared at pH 7.4 and 25°C:
No additives (solid line), LysHCl (broken line), ArgHCl (broken and dotted line), NaCl (dotted line).

- Figure 6. Viscosity of HSA solutions in the absence or presence of additives. Samples containing 305 mg/ml HSA with 200 mM amino acid or salt were prepared at pH 7.4 and 25°C.
- 353 Error bars depict the standard deviation of the mean of three independent experiments.



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