Isolation of Bovine β -Lactoglobulin from Complexes with Chitosan
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ABSTRACT

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20 A simple, economical and non-toxic method is described for the solubilization of 21 undenatured β -lactoglobulin from their complexes with chitosan. The effect of pH (8-10), 22 ionic strength (0.08-0.3M) and volume ratio between sodium acetate solutions and whey on 23 the dissociation of β-lactoglobulin-chitosan complexes was evaluated. Following a single 24 extraction step with 10 mL of 0.1M sodium acetate solution at pH 9, a recovery of 90% of 25 β-lactoglobulin with a protein purity of 95% was achieved, suggesting that electrostatic 26 interactions play a key role in the complexation of β-lactoglobulin with chitosan. The 27 presence of chitosan free in solution was ruled out according to GC-FID analysis after acid 28 hydrolysis. NMR spectroscopy showed that the β-lactoglobulin obtained and further 29 subjected to dialysis showed a structural features very similar to the native protein. This 30 method could offer a promising alternative to the traditional processes used for purification 31 of β -lactoglobulin in the dairy industry.

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34 Key words: chitosan, β -lactoglobulin, electrostatic interactions, cheese whey.

INTRODUCTION

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Bovine β -lactoglobulin (β -lg) constitutes approximately 50% of the protein present 38 39 in whey. Native β -lg is a small globular protein with defined secondary and tertiary 40 structure and is widely used as a food ingredient because of its nutritional value and its 41 ability to form gels (Allain, Paquin and Subirade, 1999; Lefevre and Subirade, 2000), 42 emulsions (Dickinson, 1997; Beaulieu, Savoie, Paquin and Subirade, 2002; Lefevre and 43 Subirade, 2003; Leung Sok Line, Remondetto and Subirade, 2005) and foams (Rantamaki, 44 Tossavainen, Outinen, Tupasela, Koskela and Kaunismaki, 2000; Bals and Kulozik, 2003). 45 Thus, much attention has been given to the development of techniques to isolate and purify 46 this protein. β -lg has been purified using various techniques including ultrafiltration 47 (Kinekawa and Kitabatake, 1996; Sannier, Bordenave and Piot, 2000), ion-exchange chromatography (Imafidon and Ng-Kwai-Hang, 1992) and precipitation or complexing 48 49 with reagents such as carboxymethyl cellulose (Hidalgo and Hansen, 1969) among others. 50 However, most of these processes have not been widely implemented for large-scale 51 purification because of their complexity, high cost, low overall yield, poor selectivity and / 52 or unacceptable product degradation associated with harsh conditions, such as high 53 temperature used during the process (Zydney, 1998; Savant and Torres, 2000; Cheang and 54 Zydney, 2003).

55 Chitosan, a polysaccharide comprising copolymers of glucosamine and *N*-56 acetylglucosamine, is positively charged at acidic pH values and interact with the 57 negatively charged polyelectrolytes such as proteins (Remuñan-López and Bodmeier, 1996; 58 Ausar et al., 2001; Casal, Corzo, Moreno and Olano, 2005). Chitosan has been used to

59 produce destabilization and coagulation of casein micelles from milk (Ausar et al., 2001), 60 reduction of protein, fat and ash contents in whey wastes (Mukhopadhyay, Talukdar, 61 Chatterjee and Guha, 2003) and to isolate caseinmacropeptide (Casal et al., 2005; Nakano, 62 Ikawa and Ozimek, 2005). Recently, we reported the complex formation between chitosan 63 and β -lg, in order to remove the β -lg from whey to simulate human milk protein (Casal, 64 Montilla, Moreno, Olano and Corzo, 2006). The same procedure could be used to recover β -lg, providing an additional step to separate the protein from the chitosan complex. 65 66 Since protein binding capacity of chitosan is affected by ionic strength and pH, the 67 aim of this work was to establish conditions for maximal extraction of β -lg from their 68 complexes with chitosan to recover high purity β -lg. 69 70 **MATERIALS AND METHODS** 71 72 Chemicals. Low-molecular-weight chitosan (LMWC), average MW 120 kDa, with 85% 73 deacetylation was supplied by Aldrich (Milwaukee, WI). Rennet powder was obtained from 74 Hansen's Laboratorium (Denmark) and starter culture Lc. Lactis ssp lactis and Lc. Lactis ssp cremoris MA0 11 from EZAL (Rhodia Iberia, Spain). Acetic acid was purchased from 75 76 Scharlau Chemie (Barcelona, Spain), chloride acid from VWR Prolabo (Fontenay sous 77 Bois, France) and sodium hydroxide from J. T. Baker (Deventer, Holland). 78 Reagents employed for GC analysis including sugar standards (glucose, galactose, 79 glucosamine), internal standard (β-phenyl-glucoside) and derivatising reagent (N-80 trimethylsilylimidazole) were obtained from Sigma (St. Louis, USA). Dried pyridine was from Merck (Darmstadt, Germany). Ultrapure water quality with 1 - 5 ppb TOC and <81

82 0.001 EU/mL pyrogen levels (Milli-Q) were produced in-house using a laboratory water
83 purification Milli-Q Synthesis A10 system (Millipore, Bellerica, Mass., USA) and was used
84 throughout.

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86 *Rennet whey preparation.* Pasteurised skimmed cow's milk inoculated with 2% of starter 87 culture was warmed to 35°C. When the pH reached 6.3, CaCl₂ (0.25 g/L), rennet powder 88 (0.1 g/L) was added. After coagulation the curd was cut to the adequate grain size 89 (approximately a rice grain), and the mixture of curd particles and whey was gently stirred 90 and heated to 45°C to reduce the moisture content in the curd. Finally, whey was filtered 91 through glass wool.

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Obtention of *B-lg-chitosan complexes*. Clarified whey obtained after treatment with 0.25 93 94 mg chitosan / mL at pH 4.5 to remove lipids and suspended solids was, then, subjected to 95 treatments with 1% (w:v) chitosan in 0.1 M acetic acid at pH 6.2 at a concentration of 1.4 96 mg chitosan / mL of whey. The mixtures were vigorously stirred for 1 min to reach a 97 complete interaction and coagulation, then allowed to stand for 30 min with gentle stirring 98 and centrifuged at 5000 g for 10 min to separate the pellets from supernatants. Under these 99 conditions, around 85% of β -lg was precipitated whereas, at least, 90% of the rest of whey proteins remained in solution (Casal et al., 2006). A control sample, containing no added 100 101 chitosan, was treated under the same conditions.

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103 *Isolation of \beta-lg from complexes with chitosan.* The pellet obtained from 1 mL of the 104 chitosan-treated whey was mixed with 5 or 10 mL of sodium acetate solutions at different 105 pHs (8, 9 or 10) and concentrations (0.08, 0.10, 0.15, 0.2 and 0.3 M), vigorously stirred for 106 30 min and centrifuged at 10000 g for 10 min to separate undissolved chitosan from 107 solubilized β -lg.

108 For total solubilization of both chitosan and β -lg, a solution of 0.1 N acetic acid at pH 4 109 was used.

110

Determination of β -lg. Analyses of whey proteins in solution were performed by RP-111 112 HPLC-UV using 0.1% (w/v) trifluoroacetic acid in double-distilled water as solvent A and 113 0.085% (w/v) trifluoroacetic acid in double-distilled water/acetonitrile (10:90, v/v) as 114 solvent B. The chromatographic conditions were those described by Casal et al. (2006). 115 Protein contents obtained in the supernatants after extraction with sodium acetate solutions 116 were expressed as a percentage of the total protein contained in the pellets. All treatments 117 were replicated three times. The identity of whey proteins was confirmed by comparison of 118 their retention times with those of the protein standards purchased from Sigma (St. Louis, 119 MO).

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121 **Determination of chitosan.** The quantitation of chitosan was done by means of their 122 glucosamine content determined by GC-FID after acid hydrolysis. The optimum hydrolysis 123 conditions that led to a high depolymerization degree of chitosan were those achieved with 124 3 N HCl at 110°C for 40 hours. Thereby, either chitosan solutions (0.01-0.2 mg/mL) or 125 supernatants containing the redissolved β -lg after extraction with 10 mL of 0.1M sodium 126 acetate at pH 9 were subjected to acid hydrolysis before subsequent GC-FID analysis.

Prior to derivatisation and GC analysis, 1 mL of filtered hydrolysates and 0.2 mL of
a solution of 0.01% (w:v) phenyl-β-D-glucoside in methanol/water (70:30, v/v) (internal

129 standard) were mixed and dried at 38-40°C in a rotavapor. Then, 100 µL of pyridine and 130 100 µL of N-trimethylsilylimidazole were added to sylilate the carbohydrates and the 131 derivatisation reaction was completed in 30 minutes at 70°C. The reaction was stopped by 132 cooling and the sylilated carbohydrates extracted with 0.1 mL of hexane and 0.2 mL of 133 water. Volumes in the range of 1-2 µL of the organic phase containing silvl derivatives 134 were injected onto the column. The trimethylsilyl ethers were separated using a commercial 135 8 m x 0.25 mm inside diameter, 0.25 µm film fused silica capillary column CP-SIL 5 CB 136 (100% dimethylsiloxane, non polar) (Chrompack, Middelburg, The Netherlands). The 137 temperatures of injector and detector were 300°C, the oven temperature was programmed 138 from 140°C to 180°C at 5°Cmin⁻¹, then at 50°Cmin⁻¹ to 250°C and keeping this temperature for 5 min, followed for an increase up to 300°C at rate of 50°Cmin⁻¹, and finally keeping 139 140 this temperature for 7 min. Injections were made in the split mode (1:10). Data acquisition 141 and integration was done using HP ChemStations software (Hewlett-Packard, Wilmington, 142 USA).

GC analysis of glucosamine gave rise to two anomers as previously shown (Karkkainen and Vihko, 1969). The quantitation of glucosamine was performed with the more retained anomer as it did not overlap with other peaks. This anomer represented 39.3% of the glucosamine. GC analyses were performed in quadruplicate. Linear correlation analyses between glucosamine and hydrolysed chitosan contents were performed using the MS Excel programme.

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150 *NMR analysis of \beta-lg.* NMR was used to compare the spectra of the chitosan-purified β -lg 151 and a commercial and native β -lg (Sigma, St. Louis, MO, USA). Prior to analysis, the

152	chitosan-purified protein was subjected to membrane filtration through an Amicon Ultra-15
153	centrifugal filter unit, 10 KDa MW cut-off (Millipore Corporation, Bedford, MA, USA),
154	followed by freeze-drying, to remove sodium acetate that may overlap with the NMR
155	protein resonances. The commercial β -lg was used directly from the container. The two
156	protein samples were dissolved in D_2O at a concentration of 20-30 mg/mL, and the pH was
157	adjusted to 2.0 with DCl.
158	Spectra were recorded in a 500 MHz Varian UNITYINOVA spectrometer (Varian
159	NMR Instruments, Palo Alto, CA, USA). All spectra were taken at 35 °C, with a spectral
160	width of 6000 Hz and a delay time of 0.5 s. The water line was suppressed using
161	presaturation for 1.5 s. One-dimension ¹ H spectra were recorded with 3.7 s acquisition time
162	and 512 transients. DQF-COSY and NOESY (100 ms mixing time) spectra were obtained
163	by accumulation of 32-64 transients, using 2048 and 512 points in the t_2 and t_1 dimensions
164	respectively. Raw data were processed and visualised using Varian software.
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166	RESULTS AND DISCUSSION
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168	Optimisation of the isolation of β -lactoglobulin from chitosan complexes
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170	Electrostatic protein-polyelectrolyte complexation sensitively depends on the
171	solution pH and ionic strength (de Vries, 2004). The pK_a of chitosan is about 6.5 (Claesson
172	and Ninham, 1992; Nyström, Kjoniksen and Iversen, 1999), thus, at pH values below 6.5
173	chitosan is positively charged along its backbone and is soluble in dilute aqueous solution,
174	whereas at higher pH dissolved chitosan usually flocculates due to deprotonation of the

amino groups (Claesson et al., 1992; Guzey and McClements, 2006). With the aim to keep chitosan undissolved and recover the β -lg free in solution, all assayed conditions were performed at pH values above the p K_a of chitosan using sodium acetate solutions as extracting solvent. Sodium acetate is a food additive recognized as safe and non-toxic by the U. S. Food and Drug Administration (FDA) and European Union.

180 Fig. 1 shows the percentage of β -lg recovered from chitosan complexes following a 181 single extraction step with 5 or 10 mL of 0.1M sodium acetate at pH 8, 9 and 10. The 182 highest recoveries were obtained extracting with 10 mL instead of 5 mL, regardless pH. Thus, around 90% of precipitated β -lg could be recovered by using 10 mL of sodium 183 184 acetate solutions at pH 9 and 10 and only 60% of the precipitated protein was resuspended 185 at pH 8, whilst hardly 35% of β -lg was extracted with 5 mL of sodium acetate solutions. 186 Additionally, a second extraction step did not increase the recovery of β -lg due to pellets 187 lost its compactness and were extremely difficult to handle.

In alkaline solution, the amino groups of the chitosan were neutralized and the β -lg remained negatively charged, leading to the lost of electrostatic linkages between the polymer and protein. The low recovery of protein observed with the sodium acetate solution at pH 8 could be due to some chitosan segments remained positively charged since the final pH of the solution was 7.0 which is pretty close to the p K_a of chitosan.

193 On the other hand, to evaluate the effect of ionic strength on the dissociation of β -lg 194 from chitosan complexes, sodium acetate solutions at pH 9 with molarities within the range 195 0.08-0.3 were assayed (Fig. 2). With 5 mL of sodium acetate solutions, only 60% of β -lg 196 was recovered at the highest molarities (0.2 and 0.3) assayed. Nevertheless, using 10 mL of 197 extracting solvent, molarities between 0.1 and 0.3 gave rise to the recovery of at least 90%

198 of the complexed protein, whereas 0.08 only allowed the dissociation of 55% of the 199 precipitated β -lg. At higher ionic strengths, the charges on both the β -lg and the chitosan 200 are shielded by the counterions (Mattison, Brittain and Dubin, 1995). As consequence of 201 these shielding effects, electrostatic interactions between chitosan and β -lg were truncated 202 allowing, thus, the presence of free β -lg in solution. Either molarities below 0.1 or reduced 203 volumes (5 mL) of sodium acetate solutions seem to be insufficient to dissociate the β -lg -204 chitosan complexes at a noticeable level.

205 Once achieved a high recovery of the precipitated β -lg, we proceed to estimate the 206 protein purity of the supernatant by RP-HPLC analysis. The supernatant comprised of 94.5% β -lg, 3% of α -lactalbumin, 2% of caseinmacropeptide and 0.3% of BSA (Fig. 3). 207 208 The minor presence of some whey proteins might be attributed to an incomplete phase 209 separation occurred during the precipitation of β -lg with chitosan and, therefore, some 210 remnants of other whey proteins could be absorbed on the pellet. Unsuccessful attempts 211 were made to obtain a better phase separation between the complexed β -lg with chitosan 212 and the rest of the whey proteins remaining in solution by increasing the centrifugation 213 speed from 5,000 to 10,000 g and/or performing several rinses of the pellet with water and 214 sodium hydroxide prior to the solubilization of β -lg (results not shown).

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216 Determination of chitosan by GC-FID following acid hydrolysis

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218 With the aim to rule out the presence of chitosan in the supernatant obtained during 219 the recovery of β -lg, a gas chromatography method after acid hydrolysis of chitosan was 220 developed for the determination of glucosamine. The graph depicted in Fig. 4A shows the 221 linear correlation between the response of glucosamine detected and levels of chitosan 222 hydrolysed. Prior to quantification of the glucosamine in the supernatants obtained during 223 the recovery of β -lg, the accuracy of the method was evaluated. Known amounts of 224 chitosan (0.0165 - 0.165 mg), within the range of chitosan added to whey for selective 225 removal of β -lg, were added to 1 mL of the supernatants and, then, subjected to acid 226 hydrolysis as described in Materials and Methods section. As shown in Fig. 4B, excellent 227 recoveries (~100%) were obtained for the whole range studied.

228 The concentration of glucosamine found in the supernatants of the solubilized β -lg 229 was 0.25 ng / mL supernatant (n=10) which is equivalent to 4 µg of chitosan per mL of 230 supernatant. To elucidate the origin of the glucosamine, i.e. whether it comes from the 231 chitosan or from whey, β -lg was precipitated by decreasing the pH with acetic acid down to 232 4.5, followed by heating at 80°C for one hour, and then was redissolved with 0.04% sodium 233 hydroxide (w:v). This sample showed a content of 0.54 ng of glucosamine / mL supernatant 234 (n=2) thus, it can be concluded that the low levels of glucosamine detected in the 235 supernatant of the resuspended β -lg were not derived from the chitosan.

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NMR analysis of the recovered β -lactoglobulin

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NMR is a powerful tool that has been used for structural analysis of milk peptides and proteins and, particularly β -lg (Belloque, 2006). Modifications in the spectral pattern indicate structural changes such as misfolding, denaturation or aggregation. To inspect the structural integrity of chitosan-purified β -lg, NMR spectra of the isolated and dialysed protein were taken. These spectra were then compared to spectra of the native protein from

244	a commercial source. The 1D and COSY spectra of the chitosan-purified and the native β -
245	lg were very similar, regarding intensity and chemical shift of resonances (results not
246	shown). This indicates that the chitosan-purified protein presented a similar structure to that
247	of the native β -lg. In addition, no line broadening was observed, consistent with a non-
248	aggregated protein. Finally, NOESY spectra cross-peaks, that represent the "structural
249	fingerprint" of the protein, were also very similar to the native protein (Fig. 5). It has to be
250	considered that the isolated protein was not tested under the same conditions that it was
251	recovered, i. e. 0.1M sodium acetate. However, if lowering the pH and removing the salts
252	led to a protein with a structure very close to that of native β -lg; it can be deduced that the
253	structure of the sodium acetate extracted protein was also able to reverse any
254	conformational changes induced by the alkaline pH or the presence of salts. This also opens
255	up the possibility of obtaining a native-like β -lg by removal of solvent co-solutes.

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In summary, from these results it can be inferred the key role of electrostatic interactions on the complexation of β -lg and chitosan. This can explain the easy solubilization, and consequently high yield of β -lg just by a simple extraction step with a solution of sodium acetate. To conclude, these findings point out that the precipitation of β lg with chitosan and subsequent solubilization may be an economical and feasible method for the recovery of β -lg present in whey. Moreover, the reversibility of the β -lg precipitation with chitosan could open up the possibility of re-using the chitosan, which

CONCLUSIONS

266	would undoubtedly make more economical the industrial application of this method to
267	purify native β -lg.
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376	FIGURE CAPTIONS

Figure 1. β-lactoglobulin (β-lg) solubilized from chitosan complexes following a single extraction step with 5 or 10 mL of 0.1M sodium acetate solution at pH 8, 9 and 10,

380 respectively. β -lg was analysed by RP-HPLC following the method described in Materials

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and Methods section. Vertical bars represent standard deviation values (n=3).
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- **Figure 2.** β -lactoglobulin (β -lg) solubilized from chitosan complexes following a single extraction step with 5 or 10 mL of 0.08, 0.1, 0.15, 0.2 and 0.3M sodium acetate solutions at
- pH 9. β-lg was analysed by RP-HPLC following the method described in Materials and Methods section. Vertical bars represent standard deviation values (n=3).
- 387
- **Figure 3.** RP-HPLC profile of the β -lactoglobulin (β -lg) solubilized from chitosan complexes following a single extraction step with 10 mL of 0.1M sodium acetate solution, pH=9. α -LA: α -lactalbumin; CMP: caseinomacropeptide.
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392 **Figure 4.** Linear correlation between (A) chitosan hydrolyzed with 3N HCl and 393 glucosamine determined by GC-FID, and (B) known amounts of standard chitosan added to 394 the supernatant containing the solubilized β -lactoglobulin and chitosan concentration 395 determined by GC-FID.

396

- 397 Figure 5. Partial view of NOESY spectra (100 ms mixing time) of (A) chitosan purified β -
- 398 lactoglobulin (β -lg) and (B) commercial β -lg in D₂O, at pH 2 and 35 °C. For more details,
- 399 see Materials and Methods section.





405 Figure 2. Montilla et al.







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Figure 5. Montilla et al.

