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# 1 Effect of alginate size, mannuronic/guluronic acid content and pH on particle 2 size, thermodynamics and composition of complexes with $\beta$ -lactoglobulin 3

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## 13 Abstract

14 Alginate is an anionic polysaccharide capable of forming insoluble particles with proteins. Hence,  
15 alginate has potential as a protein carrier. However, the role of physical properties of the polysaccharide,  
16 such as degree of polymerization ( $DP_n$ ) and mannuronic/guluronic acid ratio, remains to be fully explored.  
17 Particle formation of a high and a low molar mass alginate (ALG) with  $\beta$ -lactoglobulin (BLG) at pH 2–8  
18 depends on the average  $DP_n$  (HMW-ALG:  $1.59 \cdot 10^3$ ; LMW-ALG:  $0.23 \cdot 10^3$ ) and the mannuronic/guluronic  
19 acid ratio (1.0; 0.6) as supported by using ManA<sub>6</sub> and GulA<sub>6</sub> as models. Dynamic light scattering (DLS)  
20 showed that particles of BLG with either of the two ALGs have essentially the same hydrodynamic  
21 diameter ( $D_H$ ) at pH 3 and 2, while at pH 4 particles of LMW-ALG/BLG have larger  $D_H$  than of HMW-  
22 ALG/BLG. At pH 5–8 no significant particle formation was observed. ManA<sub>6</sub> did not form insoluble  
23 particles at pH 2–8, while GulA<sub>6</sub> formed insoluble particles, albeit only at pH 4.  $K_d$  was approximately 10-  
24 fold higher for LMW-ALG/BLG than HMW-ALG/BLG and 3 orders of magnitude higher for an alginate  
25 trisaccharide/BLG complexation as determined by isothermal titration calorimetry (ITC). The alginate  
26 trisaccharide did not form insoluble particles with BLG at pH 3 and 4, though interaction still occurred.  
27  $\Delta H_{app}$  and molar stoichiometry of BLG in the complexes with the two ALGs differed by a factor of 7, as did  
28 their  $DP_n$ , which thus affected the interaction strength, but not the BLG content. At pH 4 the BLG content  
29 doubled in the particle due to BLG dimerization. The findings emphasize the importance of  $DP_n$ ,  
30 mannuronic/guluronic acid ratio and pH in formulations containing alginate/whey protein particles.  
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36 Alginate (ALG), a linear anionic polysaccharide and major component of the cell walls of brown algae,  
37 consists of the 1,4-linked C5 epimers,  $\beta$ -D-mannuronic acid (M or ManA) and  $\alpha$ -L-guluronic acid (G or  
38 GulA) found in homo- or mixed blocks (Fig. 1) (Ci et al., 1999; Johnson, Craig, Mercer, & Chauhan, 1997;  
39 Morris, Rees, & Thom, 1980). ALG is extensively used in the pharmaceutical and food industries as a  
40 gelling and stabilizing agent (Ci et al., 1999; Johnson et al., 1997). Growing interest in nano- and micro-  
41 particles of food ingredients motivated investigations on complex formation between negatively charged  
42 polysaccharides and positively charged proteins (Aberkane, Jasniewski, Gaiani, Scher, & Sanchez, 2010;  
43 Du, Dubin, Hoagland, & Sun, 2014; Fuenzalida et al., 2016; Girard, Turgeon, & Gauthier, 2003a; Jones,  
44 Adamcik, Handschin, Bolisetty, & Mezzenga, 2010). Two types of phase separation can occur when  
45 mixing such components; repulsive phase separation known as thermodynamic incompatibility; and  
46 attractive phase separation known as thermodynamic compatibility (Doublier, Garnier, Renard, &  
47 Sanchez, 2000). The former takes place at high concentrations of neutral or similarly charged protein and  
48 polysaccharide and results in separation into a protein-rich and a polysaccharide-rich phase. By contrast,  
49 thermodynamic compatibility occurs when polysaccharide and protein carry opposite charge, polarity or  
50 similar hydrophobicity and separate into a protein/polysaccharide rich and a protein/polysaccharide poor  
51 phase or exist as one homogeneous phase (Doublier et al., 2000).  $\beta$ -lactoglobulin (BLG) is the  
52 predominant protein in whey, the major by-product of cheese making. ALG/BLG complexes can act as  
53 carriers for nutrients and nutraceuticals, e.g. folic acid and curcumin and increase colloidal stability  
54 during storage in aqueous solution (Hosseini, Emam-Djomeh, Sabatino, & Van der Meeren, 2015). ALG  
55 previously received interest as a protein carrier due to its ability to form stable particles with bovine  
56 serum albumin that maintained its conformational integrity as assessed by circular dichroism analysis  
57 after pH-induced dissociation of the particles above the protein pI (Zhao, Li, Carvajal, & Harris, 2009).  
58 The pH range suitable for formation of particles from oppositely charged polymer and protein molecules  
59 depends on protein pI and ionic strength, but not on mixing ratio and molar mass of the polysaccharide  
60 (Girard, Turgeon, & Gauthier, 2002; Weinbreck, de Vries, Schrooyen, & de Kruif, 2003). However,  
61 properties of polysaccharides in interaction with proteins such as protein affinity, complex size and  
62 amount of bound protein may depend on the DP<sub>n</sub> (Hosseini et al., 2013; Wang, Kimura, Dubin, & Jaeger,  
63 2000). If the polysaccharide is too short attractive phase separation will not occur (Li, Xia, & Dubin, 1994).  
64 The charge density of the protein is important for complexation (Kayitmazer, Seyrek, Dubin, &  
65 Staggemeier, 2003), and the charge density of the protein polymer binding site plays a key role for the  
66 strength of interaction (Comert, Malanowski, Azarikia, & Dubin, 2016). The chain flexibility of  
67 polyelectrolytes and effective charge density may also influence particle formation (Du et al., 2014;  
68 Kayitmazer, Koksal, & Iyilik, 2015). The charge density on alginate M blocks has recently been  
69 hypothesized to be less than that of G blocks (Hecht & Srebnik, 2016) which could also have an effect on  
70 particle formation (Fuenzalida et al., 2016). Here it is hypothesized that the average chain length, as well  
71 as M/G ratio of ALG affect parameters governing particle formation with BLG, such as strength of  
72 interaction and molar stoichiometry of BLG and ALG and thereby the particle size. The effect of ALG DP<sub>n</sub>  
73 and M/G-ratio on interaction with BLG was monitored by dynamic light scattering, turbidity and  
74 isothermal titration calorimetry at pH 2–9.

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## 76 2. Materials and methods

77

### 78 2.1 Materials

79

80 BLG isoform A was purified in-house from raw milk (Kristiansen, Otte, Ipsen, & Qvist, 1998) and found to  
81 be > 95 % pure as assessed by SDS-PAGE. High (HMW) and low (LMW) molar mass sodium alginates  
82 (ALGs) were produced by DuPont Nutrition and Health. HMW-ALG has number weighted average  
83 molecular mass ( $\overline{M}_n$ ) = 280 kDa, M/G ratio = 1.0 and polydispersity = 1.2. LMW-ALG has  $\overline{M}_n$  = 40 kDa, M/G  
84 ratio = 0.6 and polydispersity = 2.6 as analyzed (DuPont A/S) by SEC coupled to Multi Angle Light  
85 Scattering ( $\overline{M}_n$  and polydispersity) and FTIR spectroscopy (M/G ratio). Sodium salts of hexa-mannuronic  
86 acid (ManA<sub>6</sub>) and hexa-guluronic acid (GulA<sub>6</sub>) were purchased from Carbosynth (United Kingdom). All  
87 buffer components were of analytical grade.

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## 2.2. Methods

### 2.2.1 Sample preparation

BLG in milliQ water (270  $\mu\text{M}$ ) was prepared by stirring (150 rpm, overnight, room temperature (RT)), centrifuged (12,000 g, 20 min, 20 °C) and filtrated (0.45  $\mu\text{m}$  filter; Frisnette ApS, Denmark); the concentration was determined spectrophotometrically at 280 nm using a molar extinction coefficient  $\epsilon = 17,600 \text{ M}^{-1}\text{cm}^{-1}$  (Collini, D'Alfonso, & Baldini, 2000). HMW- and LMW-ALG (2 mg  $\text{mL}^{-1}$ ) were prepared in milliQ water by stirring (150 rpm, overnight, RT) to ensure complete dispersion and filtrated (0.45  $\mu\text{m}$  filter; Frisnette ApS, Denmark). GulA<sub>6</sub> and ManA<sub>6</sub> (104 mM) were dissolved in milliQ water overnight and centrifuged (as above). BLG and ALG stocks were diluted with buffers (stock 100 mM, final 10 mM): glycine (pH 2), sodium citrate/citric acid (pH 3–6) and Tris-HCl (pH 7–9) and mixed to the desired ratios. For ITC BLG was dissolved in the respective buffers (200  $\mu\text{M}$ ), centrifuged (20,000 g, 20 min, 20 °C) and dialysed against 10 mM sodium citrate/citric acid (3 x 4 h, 4 °C, 6–8 kDa cutoff, SpectraPor membrane; Spectrum). ALGs (4 mg  $\text{mL}^{-1}$ ) were dissolved in 10 mM sodium citrate/citric acid pH 3 or 4, centrifuged and dialysed (as above) to remove sodium ions added with the ALG. ALG tri-saccharide (ALGOS; see section 2.2.2) was dissolved in water (8.4 mM) and 100 mM sodium citrate/citric acid pH 3 or 4 was added to a final buffer concentration of 10 mM followed by centrifugation (as above).

### 2.2.2 Production, purification and characterization of ALGOS

LMW-ALG (10 mg  $\text{mL}^{-1}$ ) in 50 mM Tris pH 7.2, 1 mg  $\text{mL}^{-1}$  BSA (100 mL) was added 10 U  $\text{mL}^{-1}$  of an endoacting alginate lyase from *Sphingomonas sp.* (Megazyme, United Kingdom) and incubated for 6.5 h at 40°C under gentle mixing. The ALGOS obtained was desalted in milliQ water (HiPrep Desalt 26/10; GE Healthcare, Denmark) and purified by anion exchange chromatography (Mono Q 5/50 GL; GE Healthcare, Denmark) eluted by a 0–2 M NaCl linear gradient (2 h) in milliQ water at a flow rate of 1  $\text{mL min}^{-1}$ . Eluted ALGOS was monitored by the absorbance at 235 nm (Park, Kam, Lee, & Kim, 2012). Fractions were pooled, added 0.5 M NaCl, filtered (3 kDa cut-off centrifugal filter; Amicon, Germany) to remove remaining protein, desalted as above and freeze dried (ScanVac FreezeSafe Freeze Dryer, Holm & Halby, Denmark). ALGOS samples were subjected to TLC (aluminium sheet Silica gel 60 WF254; Merck, Germany) developed twice in butanol:acetic acid:QH<sub>2</sub>O (2:1:1) and stained by 10 % sulfuric acid, 80 % ethanol, 8 % H<sub>2</sub>O and 2 % orcinol at 300°C. MALDI-TOF MS was performed (Ultraflex II TOF/TOF; Bruker Daltonics) in positive ion linear mode using a polished steel TF MTP 384 target (Bruker Daltonics GmbH). Peak analysis of mass spectra was done using FlexAnalysis Version 3.3 (Bruker Daltonics GmbH).

### 2.2.3 Determination of alginate concentration and preparation of alginate standard

LMW-ALG (3 mg  $\text{mL}^{-1}$ ) was dissolved in milliQ water, centrifuged and the supernatant dialyzed as above (3 x 4 h) against milliQ water. A 25 mL volumetric flask with glass stopper was cleaned with 96 % ethanol, heated (80 °C, 1 h), cooled to RT, weighed, filled with ALG solution, freeze-dried, desiccated (36 h, vacuum, RT; ME1 pump, Vacuubrand, United Kingdom), weighed, filled to the 25 mL mark with 10 mM sodium citrate/citric acid pH 4, and the LMW-ALG was dissolved under magnetic stirring (12 h) to give an ALG standard solution. The concentration of ALG used for ITC was determined by the phenol sulfuric acid method (Dubois, Gilles, Hamilton, Rebers, & Smith, 1956) with this ALG standard.

### 2.2.4 Turbidimetric and UV absorbance measurements

Turbidity of ALG/BLG mixtures (0.2 mg  $\text{mL}^{-1}$ /54  $\mu\text{M}$ ) was measured spectrophotometrically at 600 nm (Ultrospec pro 2100; Amersham Biosciences). Samples were centrifuged (20,000 g, 20 min, 20 °C; Eppendorf centrifuge 5417R, Denmark) and the absorbance of supernatants measured at 280 nm.

### 2.2.5 Dynamic light scattering

Particle size distribution of ALG (1 mg  $\text{mL}^{-1}$ ), BLG (54  $\mu\text{M}$ ) and ALG/BLG (0.2 mg  $\text{mL}^{-1}$ /54  $\mu\text{M}$ ) mixtures was analysed by DLS (DLS instrument BI-200SM Brookhaven Instruments Corporation; USA) at a scattering angle of 90° at 23 °C. The distributions of mean apparent translational diffusion coefficients

146 ( $D_T$ ) were determined by fitting DLS autocorrelation functions obtained with the instrument using non-  
147 negative constrained least-squares (NNLS). The distribution of mean apparent  $D_T$  was converted to the  
148 distribution of hydrodynamic diameter ( $D_H$ ) using the Stokes-Einstein equation:

$$D_H = kT/3\pi\eta D_T$$

152 where  $k$  is the Boltzmann constant,  $T$  the absolute temperature, and  $\eta$  the solvent viscosity (0.93 mPa x s;  
153 assumed to be that of water at 296 K).

### 155 2.2.6 $\zeta$ -Potential of alginate and $\beta$ -lactoglobulin

157 Electrophoretic mobility measurements were performed (Brookhaven 90Plus ZetaPALS Potential  
158 Analyzer; Brookhaven Instruments Corporation; USA) and the zeta potential (represented by  $\zeta$  in  
159 millivolts) of BLG (1 mg·mL<sup>-1</sup>) and LMW-ALG (1 mg·mL<sup>-1</sup>), HMW-ALG (1 mg·mL<sup>-1</sup>) was obtained from the  
160 electrophoretic mobility ( $\mu_e$ ) using Helmholtz-Smoluchowski equation:

$$\mu_e = \varepsilon\zeta/\eta$$

164 where  $\varepsilon$  is the dielectric constant (water) multiplied by the permittivity of vacuum, and  $\eta$  the solvent  
165 viscosity (0.93 mPa x s; assumed to be that of water at 296 K).

### 167 2.2.7 Isothermal titration calorimetry

169 ITC was used to determine the apparent dissociation constant ( $K_d$ ) and the apparent change in  
170 enthalpy ( $\Delta H_{app}$ ) for complexation of ALGs and ALGOS with BLG at pH 3 and 4. Titrations were done with  
171 dialyzed ALG (1.5–3.5 mg mL<sup>-1</sup>; 5–88  $\mu$ M, diluted in dialysis buffer) in the syringe and BLG (50–80  $\mu$ M,  
172 diluted in dialysis buffer) in the cell, by 26–40 injections each of 6  $\mu$ L (NanoITC2G; TA Instruments, USA)  
173 including an initial 3  $\mu$ L injection (deleted from the data set). A blank titration adding ALG into buffer was  
174 subtracted as heat of dilution. The conventional binding model for  $n$  identical sites was fitted to the  
175 resulting data (NanoAnalyze Data Analysis Version 3.4.0; TA instruments, USA). It is noted that this model  
176 relies on the mass action description of the binding process, which is only an approximation for the  
177 current system as the binding of BLG will modify the surface potential of ALG, hence resulting in  
178 continuous reduction of the affinity for BLG.  $K_d$  is thus not a true dissociation constant but an empirical  
179 measure of average affinity.  $K_d$  and  $\Delta H_{app}$  were calculated by the program (NanoAnalyze software, TA).  
180 Errors are standard deviations for the regression analysis (NanoAnalyze software). This setup ensures  
181 similar ionic strength at all pH as opposed to particle formation by acidifying protein ALG mixtures.  
182 Purified ALGOS (in syringe: 4 mg mL<sup>-1</sup> = 7.6 mM) was added in 20 injections into BLG (in cell: 500  $\mu$ M),  
183 each of 2  $\mu$ L including an initial 0.4  $\mu$ L injection (excluded from the data set) (ITC200; Thermo Scientific,  
184 USA). Thermograms were corrected for heat of dilution, obtained from a blank titration, and a one-site  
185 binding model was fitted to integrated and normalized binding data.

## 187 3. RESULTS AND DISCUSSION

188

### 189 3.1 $\zeta$ -potential of ALG and BLG

191 The  $\zeta$ -potential was measured in order to determine the surface charge under conditions where  
192 particles are formed.  $\zeta$ -potential is a measure of the electrokinetic potential, *i.e.* the difference in electric  
193 potential between the solvent and the stationary layer of solvent molecules bound to dissolved particles.  
194 A  $\zeta$ -potential is relative to the surface charge of molecules in solution (Makino & Ohshima, 2010). HMW-  
195 and LMW-ALGs have  $\overline{M}_n$  of 280 kDa and 40 kDa with calculated average maximum number of negative  
196 charges at high pH of 1591 and 227 as derived from  $M$  and  $G$  of 194 Da and  $pK_a$  3.38 and 3.65,  
197 respectively (Draget, Braek, & Smidrod, 1994). At pH 2 HMW- and LMW-ALGs have a  $\zeta$ -potential of -36  
198 and -19 mV, respectively, indicating a negative surface charge even at this low pH (Fig. 2), perhaps due to  
199 polycarboxylic acid effects on  $pK_a$  and reflecting inter- and intra-molecular hydrogen bonding and

200 electrostatic forces (Castaneda et al., 2009). The  $\zeta$ -potential did not change at pH 4. The different pH  
201 required to change the  $\zeta$ -potential of HMW-ALG (pH 5) and LMW-ALG (pH 6) may be due to their  
202 different M/G ratio. BLG displayed positive  $\zeta$ -potential of +24 mV at pH 2, which decreased linearly with  
203 increasing pH to -19 mV at pH 9 and a  $\zeta$ -potential of 0 mV being reached just below pH 5 in accordance  
204 with experimental and calculated (ProtParam) BLG pI values of 4.7–5.2 (Bromley, Krebs, & Donald, 2005;  
205 Das & Kinsella, 1989; Sawyer & Kontopidis, 2000) and 4.83, respectively. The data agree with the  $\zeta$ -  
206 potential value reported (Harnsilawat, Pongsawatmanit, & McClements, 2006) to be 0 mV for BLG at pH 5  
207 and negative for ALG of similar mass to HMW (216 kDa), respectively, in the same pH range as studied  
208 here. Overall BLG and ALG carry opposite charge below pH 5. Notably, the  $\zeta$ -potential of HMW- and  
209 LMW-ALGs differs by a factor of 3, which is less than expected from the  $DP_n$  of the ALGs varying 7 fold. An  
210 explanation for this may be the difference in viscosity.

211

### 212 3.2 Turbidity and absorbance of ALG/BLG mixtures

213

214 The solubility of ALG/BLG complexes was assessed by turbidity (at 600 nm) and absorbance (at 280  
215 nm) measurements. The turbidity increased below pH 5 and did not change above pH 5, indicating  
216 attractive electrostatic interactions of ALG and BLG, which carry opposite charge at pH < 5 (Fig. 2 and 3A).  
217 No increase in turbidity or decrease in absorbance was observed for samples containing BLG and no ALG  
218 (data not shown). Decrease in turbidity was observed for all samples at pH 2 relative to pH 4, suggesting  
219 dissociation of particles as confirmed by an increased amount of protein in solution after centrifugation.  
220 At pH 4 and 3 a large decrease of protein in solution indicated formation of insoluble ALG/BLG particles.  
221 The reported optimum at pH 3.65–3.8 for formation of ALG/BLG particles with ALG of similar molar mass  
222 (200 kDa) (Hosseini et al., 2013) agrees with the present finding of highest turbidity of ALG/BLG at pH 4  
223 (Fig. 3A).

224 Addition of ManA<sub>6</sub> or Gula<sub>6</sub> to BLG resulted in negligible turbidity at pH 2–9, except at pH 4, where Gula<sub>6</sub>  
225 elicited higher turbidity than ManA<sub>6</sub>, accompanied by significant decrease in the amount of soluble  
226 protein measured at 280 nm after centrifugation (Fig. 3B). No drop in soluble protein was observed at pH  
227 4 for ManA<sub>6</sub>/BLG. This suggests that ALG/BLG particle formation is driven by G rather than M, possibly  
228 reminiscent of the greater ability of Gula<sub>6</sub> to form particles with BLG (Fig. 3C,D), although relatively  
229 higher Gula<sub>6</sub> concentration was required compared to LMW- and HMW-ALG (supplementary Fig. S2). The  
230 same concentration of ManA<sub>6</sub> marginally increased turbidity without significant loss of soluble protein.  
231 Noticeably, even the hexa-saccharide can participate in insoluble particle formation with BLG. The  
232 greater ability of Gula<sub>6</sub> to form insoluble complexes than ManA<sub>6</sub> was confined to a narrow pH range  
233 compared to the ability of HMW- and LMW-ALG (Fig. 3A,B). ALGOS/BLG mixtures did not display any  
234 turbidity at pH 3 and 4 and remained in solution after centrifugation ( $A_{280}$  of ALGOS/BLG was 0.941  
235 before and 0.934 after centrifugation). Poly-G blocks are known to form a “buckled”-chain conformation  
236 with higher affinity for calcium than other metal cations, whereas M rich regions or mixed M/G blocks do  
237 not discriminate between different metal cations (Wong, Preston, & Schiller, 2000). The charge density  
238 simply may be higher on the more rigid Gula<sub>6</sub> (Hecht & Srebnik, 2016) than on ManA<sub>6</sub>. Gula<sub>6</sub> therefore is  
239 superior in neutralizing local surface charge on BLG leading to aggregates as when the overall charge is 0  
240 at the protein pI. This is in line with the observation that polyelectrolyte flexibility and local charge  
241 density is important for binding (Du et al., 2014). ALG of 7 kDa and M/G ratio 5 was recently reported to  
242 bind lysozyme more weakly than ALG of 4 kDa and M/G ratio 1.42 (Fuenzalida et al., 2016). A similar  
243 effect of G blocks as seen with calcium/ALG and lysozyme/ALG interactions possibly occurs for ALG  
244 binding BLG.

245

### 246 3.3 Size of HMW- and LMW-ALGs/BLG particles

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248 The average particle size distribution of ALG, BLG and their mixtures at pH 2–8 showed a maximum at  
249 pH 4 for ALG/BLG, as given by the hydrodynamic diameter ( $D_H$ ) (Fig. 4). Lower  $D_H$  values at pH 3 and 2  
250 were similar for particles of HMW-ALG/BLG and LMW-ALG/BLG. The particle size at pH 3 ( $1482 \pm 93$  nm  
251 and  $1652 \pm 92$  nm for LMW and HMW respectively) is slightly smaller than observed previously  
252 (Qomarudin et al., 2015), which may be due to differences in ionic strength of the buffer. The particle  
253 formation begins at pH 5, where  $D_H$  increases slightly compared to the BLG control.  $D_H$  of ALG/BLG  
254 particles were smaller at pH 3 and 2 than pH 4, suggesting partial dissociation of particles at low pH as  
255 reflected by higher  $K_d$  and lower stoichiometry of ALG/BLG at pH 3 (Table 1).  $D_H$  is highest for LMW-ALG  
256 compared to HMW-ALG at pH 4 and 5 which may be due to the higher G content of LMW-ALG in  
257 agreement with formation of more insoluble complexes with BLG by Gula<sub>6</sub> than by ManA<sub>6</sub> at pH 4 (Fig.

258 3B,C).  $D_H$  increased marginally at pH 6–8 compared to the BLG control (likely due to the  $D_H$  being  
259 calculated as a single species) indicating no interaction and supporting that the interaction is ionic, as BLG  
260 and ALG both carry a negative net charge at pH 6–8 (Fig. 2). The HMW- and LMW-ALG controls displayed  
261 almost the same  $D_H$  at pH 2–8 with a slight increase at pH 8 (Fig. 4, bottom). This may stem from greater  
262 charge at higher pH and hence stronger repulsion between individual saccharide units/regions.  
263

### 264 3.5 Isothermal titration calorimetry

265  
266 The large differences in particle size of ALG/BLG complexes between pH 4 and 3 may reflect variation  
267 in affinity and molecular stoichiometry. At pH 3 HMW-ALG/BLG complex formation has 11-fold lower  $K_d$   
268 of  $23 \pm 3$  nM than  $K_d$  of  $266 \pm 48$  nM with LMW-ALG, different from what is expected from the avidity  
269 effect, possibly due to the difference in M/G ratio. HMW-ALG binds 7.6 times more BLG than LMW-ALG,  
270 corresponding to their 7-fold difference in  $\overline{M}_n$  (Table 1). At pH 4 both the strength of interaction and the  
271 stoichiometry increased two-fold in accordance with the larger  $D_H$  compared to at pH 3 (Fig. 4). HMW-  
272 and LMW-ALGs both displayed exothermic  $\Delta H_{app}$  of complex formation with BLG at pH 3 and 4 (Fig. 5).  
273 The ratio of  $\Delta H_{app}$  of particle formation at pH 3 between HMW- and LMW-ALGs was 6.7 ( $-6271.0 \pm 96.6$   
274  $\text{kJ mol}^{-1}$ ,  $-939.5 \pm 27.4 \text{ kJ mol}^{-1}$ ) and at pH 4 6.6 ( $-6666.0 \pm 125.0 \text{ kJ mol}^{-1}$ ,  $-1013.0 \pm 24.0 \text{ kJ mol}^{-1}$ ) (Table  
275 1) but when compared in terms of  $\text{J g}^{-1}$  none of the observed enthalpy's are significantly different.

276  $\Delta H_{app}$  of binding is similar to that of complex formation reported for BLG and other polysaccharides. Thus  
277  $\Delta H_{app}$  was  $-75.7 \text{ kJ (mol galacturonic acid)}^{-1}$  for pectin/BLG interaction (Girard, Turgeon, & Gauthier,  
278 2003b) and very large  $\Delta H_{app}$  was found in related systems undergoing attractive phase separation  
279 (Aberkane et al., 2010; de Souza, Bai, Goncalves, & Bastos, 2009). When soluble protein and anionic  
280 polysaccharides interact below pI of the protein, particle size increases significantly (Fig. 4). This is  
281 accompanied by reduction of the solvent accessible surface area and hence the number of hydrogen  
282 bonds with solvent, which would add an enthalpy effect. If this, however, was the sole explanation for  
283  $\Delta H_{app}$ , the reaction, due to loss of hydrogen bonds, would be endothermic which was not the case (Fig. 5).  
284 Another effect on  $\Delta H_{app}$  apart from the change in enthalpy of interaction is the amount of bound protein  
285 that leaves solution per molecule of ALG, which at both pH 3 and 4 are similar when expressed in terms  
286 of  $\text{J g}^{-1}$ . The strong exothermic signal could be the result of bringing separated opposite charges together,  
287 which may also explain that  $\Delta H_{app}$  does not differ much between pH 3 and 4 in accordance with the  $\zeta$ -  
288 potential of the ALGs not changing significantly in that pH range (Fig. 2).

289 Since ALGOS did not display any turbidity at pH 3 and 4 the interaction with BLG was confirmed by using  
290 ITC. ALGOS and BLG interact at pH 4 with  $K_d = 568 \pm 40 \mu\text{M}$  and  $\Delta H_{app} = -3.4 \pm 0.5 \text{ kJ mol}^{-1}$  (Table 1) and at  
291 pH 3 with  $K_d = 1028 \pm 203 \mu\text{M}$ , possibly reflecting this is closer to the  $\text{p}K_a$  of ALGOS. The different  $n$  values  
292 of 1.71 and 1.28 bound ALGOS molecules per BLG monomer at pH 3 and 4, respectively (Table 1),  
293 probably reflect the higher positive charge of BLG (Fig. 2) as well as the larger available surface area at pH  
294 3, where monomeric BLG prevails (Sakurai, Oobatake, & Goto, 2001).  $\Delta H_{app}$  seems unaffected by the pH  
295 difference as observed also with the two ALGs, but is much smaller than  $\Delta H_{app}$  for the polysaccharides  
296 (Table 1). This may be due to particle formation not occurring with ALGOS. The much higher affinity of  
297 ALG than ALGOS for BLG at pH 4 and 3 suggested massively enhanced interaction of BLG with the  
298 polysaccharide, likely as reflecting an avidity effect. This confirms previous findings for lysozyme that the  
299 larger the ALG  $\text{DP}_n$  the higher the affinity for this positively charged protein (Fuenzalida et al., 2016).  
300 Along the same line,  $\Delta H_{app}$  is much smaller for ALGOS than for ALGs. The formed ALGOS/BLG complexes  
301 at completion of the titration are soluble as assessed by turbidity and  $A_{280}$  measurements. This supports  
302 the hypothesis of Li et al. that if  $\text{DP}_n$  of a polyelectrolyte is too low, large particles are not formed with  
303 oppositely charged molecules (Li et al., 1994) even though interaction still occurs (Fig. 5C). In the present  
304 case a too low  $\text{DP}_n$  is  $< 6$  as judged from the  $\text{ManA}_6$  and  $\text{Gula}_6$  BLG interactions.

305 LMW-ALG binds  $15.0 \pm 0.3$  and  $30.5 \pm 0.5$  BLG monomers at pH 3 and 4, respectively, while HMW-ALG  
306 binds  $113.8 \pm 1.1$  and  $238.6 \pm 3.3$  BLG monomers (Table 1). This two-fold difference in stoichiometry  
307 between pH 3 and 4, may reflect ALG binding with monomeric and dimeric form of BLG (Sakurai et al.,  
308 2001; Taulier & Chalikian, 2001), and suggests that one or more ALG binding sites are situated close to  
309 the BLG dimerization interface. If a binding site on BLG was located far from the dimerization interface  
310 the amount bound per monomer should be similar at pH 3 and 4, but if the binding site is close to the  
311 dimerization interface a BLG dimer can bind only one ALG molecule. This is further supported by the  $K_d$   
312 value being 50 % at pH 4 compared to pH 3, as expected from the avidity effect of dimerization. Girard et  
313 al. previously suggested BLG residues 132–148 contain a pectin binding site (Girard et al., 2003a). When  
314 viewed in the quaternary structure of the BLG dimer (Brownlow et al., 1997), this region forms a cleft  
315 possibly functioning as a binding site located in the dimerization interface. Altogether, the particle size of

316 ALG/BLG particles is greatly affected by pH but unaffected by DP<sub>n</sub> in the range pH 2–3. At pH 4 G-blocks  
317 have a greater tendency to form insoluble particles than M-blocks, which may explain why LMW/BLG  
318 particles are larger than HMW/BLG particles at pH 4. The range in which insoluble particles are formed  
319 also depends on DP<sub>n</sub> with hexasaccharides only forming insoluble particles at pH 4 and trisaccharides  
320 (ALGOS) never forming insoluble particles which may have something to do with the strength of  
321 interaction as seen from the ITC analysis. The present results indicate that the ALG/BLG particle  
322 formation varies importantly within the pH 2–4 range, probably due to BLG dimerization and to the local  
323 availability of charges on ALG as well as effects arising from the M/G ratio.

#### 324 4. Conclusion

325

326 DP<sub>n</sub> of ALG has a prominent effect on affinity and molar binding stoichiometry of BLG as determined  
327 by ITC, showing that the stoichiometry for the LMW- and HMW-ALGs increases by a factor corresponding  
328 to their difference in DP<sub>n</sub>. Small oligosaccharides interact with mM-affinity and do not form insoluble  
329 particles, but when binding is accumulated in case of polysaccharides, the combined interaction is in the  
330 nM-range and insoluble particles are formed. While the D<sub>H</sub> of complexes was unaffected by ALG DP at pH  
331 3 and 2, it was greatly influenced in the pH 3–5 range and insoluble ALG/BLG complexes appeared at pH 3  
332 and 4. This warrants further study using the model ALGOS and BLG to localize binding sites by identifying  
333 BLG residues engaged in complex formation. The results underline the importance of considering DP<sub>n</sub>,  
334 M/G ratio and pH when preparing formulations for dairy food products from ALG and whey proteins.

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336

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341

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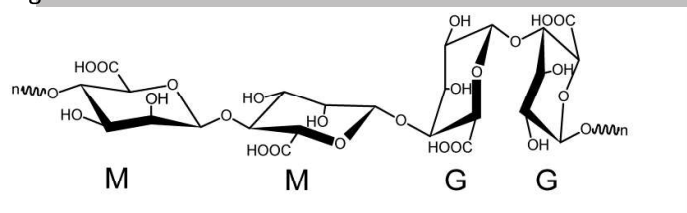
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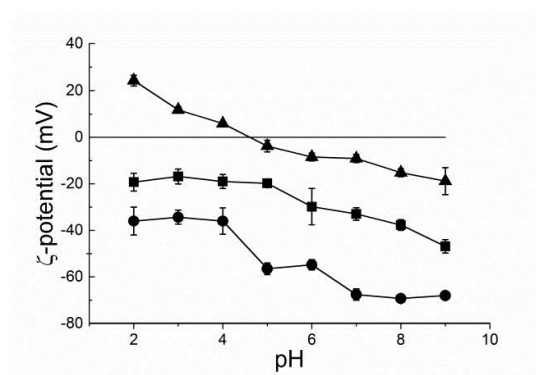


450

451 Fig 1. Chair conformation illustrating the structural motifs; 1,4-linked  $\beta$ -D-mannuronate block (left);  $\alpha$ -L-  
 452 guluronate block (right) and mixed M/G block (center) of ALG at low pH. n represents the continued  
 453 polysaccharide.

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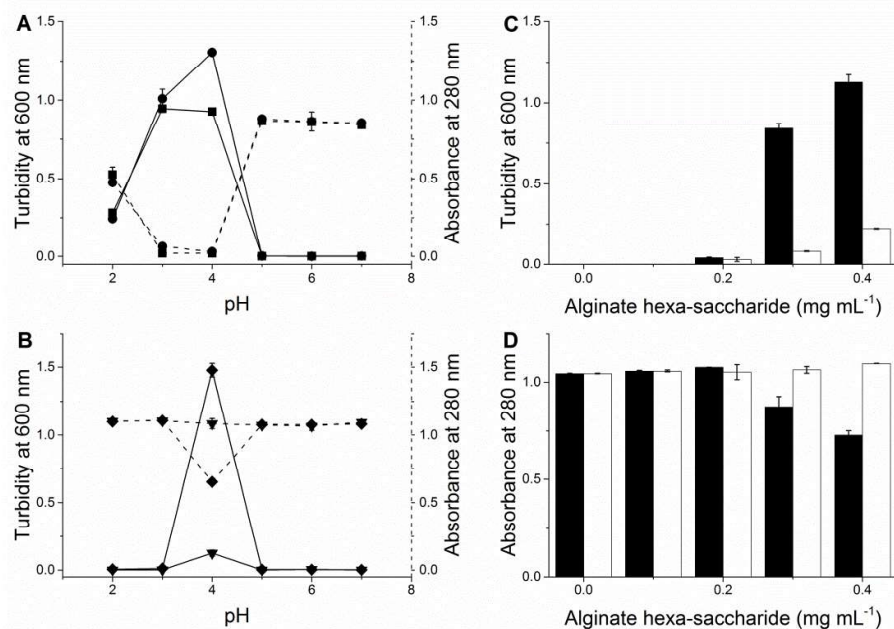
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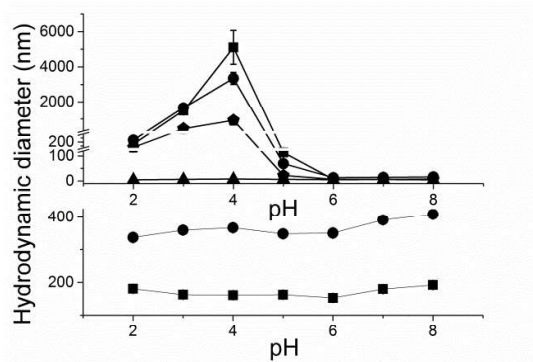
457 Fig 2.  $\zeta$ -potential for LMW-ALG, HMW-ALG and BLG as a function of pH. HMW-ALG (circle), LMW-ALG  
 458 (square), BLG (triangle).

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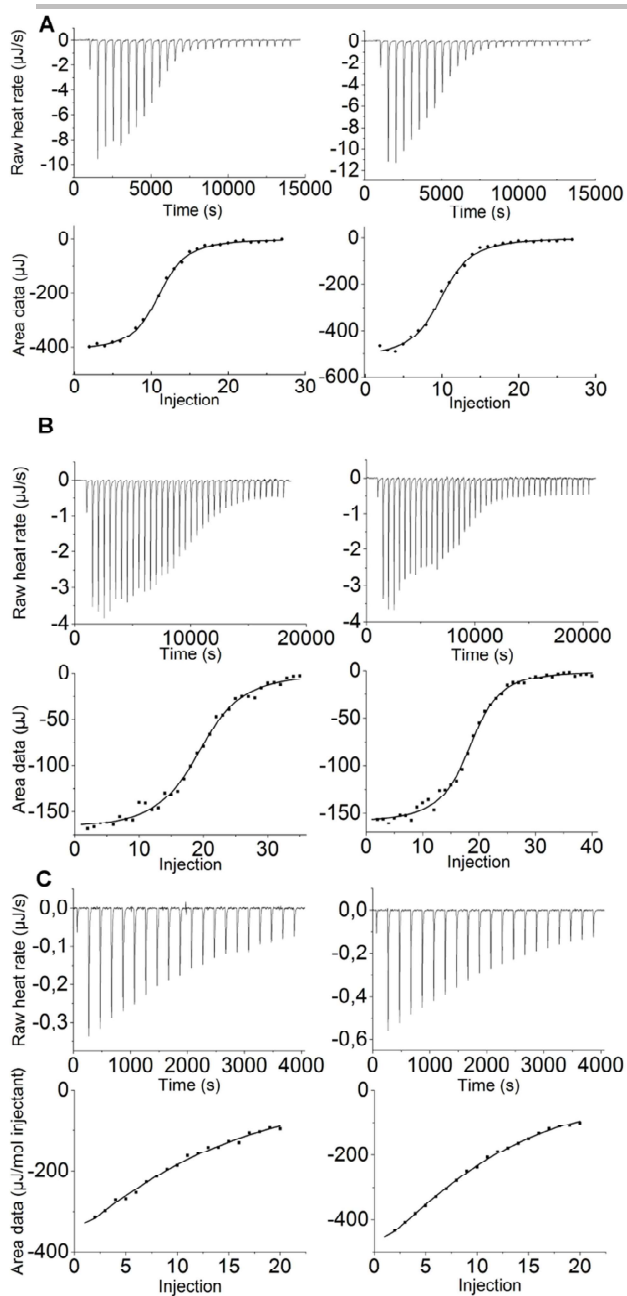


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461 Fig 3. Turbidity and absorbance of ALG/BLG mixtures as a function of pH. Turbidity was measured at 600  
 462 nm after mixing (solid line). Absorbance of the supernatant was measured at 280 nm to determine the  
 463 amount of protein remaining in solution (dashed line). A) 0.2 mg mL<sup>-1</sup> HMW- (circle) or LMW-ALG  
 464 (square) mixed with 54 μM BLG. B) 0.4 mg mL<sup>-1</sup> ManA<sub>6</sub> (diamond) or Gula<sub>6</sub> (upside down triangle) mixed  
 465 with 1 mg mL<sup>-1</sup> BLG. C) Turbidity of 1 mg mL<sup>-1</sup> BLG mixed with varying concentration of Gula<sub>6</sub> (black) or  
 466 ManA<sub>6</sub> (white) at pH 4. D) Absorbance after centrifugation of 1 mg mL<sup>-1</sup> BLG mixed with varying  
 467 concentration of Gula<sub>6</sub> (black) or ManA<sub>6</sub> (white) at pH 4.  
 468



484 Fig 4. Hydrodynamic diameter ( $D_H$ ) of ALGs, BLG and ALG/BLG mixtures. Top: Hydrodynamic diameter of  
485 ALG/BLG as a function of pH. LMW-ALG (0.7  $\mu\text{M}$ ) and BLG (54  $\mu\text{M}$ ) (square), HMW-ALG (0.1  $\mu\text{M}$ ) and BLG  
486 (54  $\mu\text{M}$ ) (circle), LMW-ALG (0.1  $\mu\text{M}$ ) and BLG (54  $\mu\text{M}$ ) (pentagon) and BLG (54  $\mu\text{M}$ ) (triangle). Bottom:  
487 Hydrodynamic diameter of ALGs (1  $\text{mg mL}^{-1}$ ) as a function of pH. LMW-ALG (square), HMW-ALG (circle).  
488



489

490 Fig 5. ITC of ALG/BLG complex formation at pH 3 and 4. For each experiment the upper portion shows the  
491 raw data with the baseline subtracted and the lower part shows the integrated peaks with the fitted one  
492 site binding model. A) HMW-ALG (left) LMW-ALG (right) titrated into BLG at pH 3. B) HMW-ALG (right)  
493 and LMW-ALG (left) titrated into BLG at pH 4. C) ALGOS titrated into BLG at pH 3 (left) and pH 4 (right).

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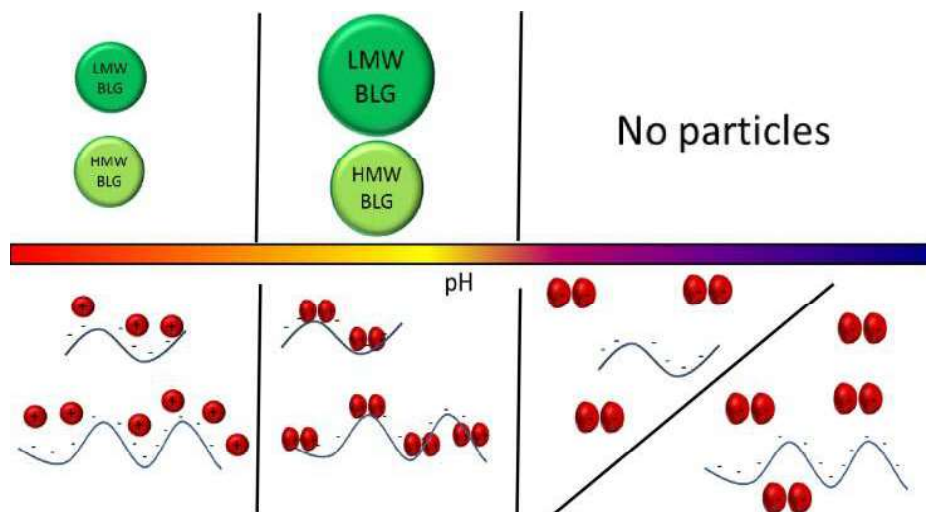
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500 Table 1. Thermodynamic parameters obtained for ALGs and BLG at pH 3 and 4 by using ITC (Fig. 5).  
 501 \*stoichiometry is reported as alginate oligomers bound per monomer of BLG. Numbers in brackets  
 502 represent the data in terms of mass.

ALG	pH	$K_d$ [nM] (g L <sup>-1</sup> )	n [BLG monomers/molecule ALG] (g BLG/g ALG)	$\Delta H_{app}$ [kJ mol <sup>-1</sup> ] (J g <sup>-1</sup> )
HMW	3	23 ± 3 (6.44 · 10 <sup>-3</sup> ± 7.46 · 10 <sup>-4</sup> )	113.8 ± 1.1 (7.44 ± 0.08)	-6271.0 ± 96.6 (-22.40 ± 0.35)
LMW	3	266 ± 48 (1.06 · 10 <sup>-2</sup> ± 1.87 · 10 <sup>-3</sup> )	15.0 ± 0.3 (6.88 ± 0.14)	-939.5 ± 27.4 (-23.41 ± 0.68)
ALGOS	3	1028 · 10 <sup>3</sup> ± 203 · 10 <sup>3</sup> (0.54 ± 0.11)	1.71 ± 0.2* (20.9 ± 0.06)	-3.1 ± 0.6 (-4.04 ± 0.40)
HMW	4	12 ± 2 (3.36 · 10 <sup>-3</sup> ± 5.14 · 10 <sup>-4</sup> )	238.6 ± 3.3 (15.59 ± 0.21)	-6666.0 ± 125.0 (-23.81 ± 0.45)
LMW	4	119 ± 22 (4.76 · 10 <sup>-3</sup> ± 8.83 · 10 <sup>-4</sup> )	30.5 ± 0.5 (13.94 ± 0.24)	-1013.0 ± 24.0 (-25.31 ± 0.62)
ALGOS	4	568 · 10 <sup>3</sup> ± 40 · 10 <sup>3</sup> (0.30 ± 0.02)	1.28 ± 0.0* (27.10 ± 0.00)	-3.4 ± 0.2 (-6.49 ± 0.33)



## Graphical abstract



**Highlights**

- Particle size of alginate/ $\beta$ -lactoglobulin complexes depends on pH
- Alginate/ $\beta$ -lactoglobulin complex size depends on alginate M/G ratio and/or the molar mass
- Poly-guluronic acid has greater nanoparticle formation tendency than poly-mannuronic acid
- Formation of nanoparticles depends on alginate DP<sub>n</sub> and pH
- With alginate oligosaccharide of DP<sub>n</sub> = 3 no particles are formed but interaction occurs