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Additional Information

## **Gums induced microstructure stability in Ca(II)-alginate beads containing lactase analyzed by SAXS**

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## **ABSTRACT**

Previous works show that the addition of trehalose and gums in  $\beta$ -galactosidase (lactase) Ca(II)-alginate encapsulation systems improved its stability against freezing and dehydration processes. However, there is no available information on how the microstructure is affected due to the constraints imposed by the operation conditions. The aim of this research is to study the microstructural changes of Ca(II)-alginate matrices driven by the presence of trehalose, arabic and guar gums as excipients and to correlate these changes with the enzymatic activity of the encapsulated lactase.

The structural modifications at different scales were assessed by SAXS. The incorporation of gums as second excipients induces a significant stabilization in the microstructure not only at the rod scale, but also in the characteristic size and density of alginate dimers (basic units of construction of rods) and the degree of interconnection of rods at a large scale, improving the performance in terms of lactase activity.

*Keywords:* alginate beads, hydrocolloids, encapsulation,  $\beta$ -galactosidase, microstructure, Small-Angle X-ray scattering (SAXS).

## 1. Introduction

In recent years, the encapsulation of  $\beta$ -galactosidase (lactase) in order to control its release and improve its stability against thermal and mechanical effects has been widely investigated (Traffano-Schiffo, Castro-Giraldez, Fito, & Santagapita, 2017a; Zhang, Zhang, Chen, McClements, & 2016; Zhang, Zhang, & McClements, 2017; Estevinho, Damas, Martins, & Rocha, 2014).

Alginate is one of the most used anionic polyelectrolytes for the encapsulation of bioactive compounds (Santagapita, Mazzobre, & Buera, 2012). It consists of (1 $\rightarrow$ 4)-linked residues of  $\beta$ -D-mannuronate (M) and  $\alpha$ -L-guluronate (G), and in solution it generates a hydrogel matrix through the complexation of G-blocks with di- or trivalent cations such as  $\text{Ca}^{2+}$ ,  $\text{Mn}^{2+}$ ,  $\text{Zn}^{2+}$ ,  $\text{Al}^{3+}$ ,  $\text{Cr}^{3+}$ , and  $\text{Ce}^{3+}$  (Santagapita, Mazzobre, & Buera, 2011), embodying them into the cavities formed by a cooperative pairing of contiguous G-blocks (Stokke et al., 2000) forming the structure commonly known as “egg-box” (He, Liu, Li, & Li, 2016). Additionally, the sodium alginate solution can be gelled at acidic pH. Once a critical fraction of carboxylate residues are protonated, the decrease in the polymer charge density allows for chain-chain interactions leading to gelation. The inherent pH for the consolidation of H–alginate hydrogel is around 3.5, depending on the mannuronic ( $\text{p}K_{\text{a}} = 3.38$ ) to guluronic acid ( $\text{p}K_{\text{a}} = 3.65$ ) relative content of the employed alginate. Moreover, the Ca(II)–alginate structure can be obtained by cation exchange from a parent H–alginate hydrogel (Sonego, Santagapita, Perullini, & Jobbágy, 2016), indicating a higher affinity of alginate polymer for  $\text{Ca}^{2+}$ . It has been demonstrated that the combined use of alginate with sugars and/or other biopolymers such as

arabic and guar gums allows to improve the stability of enzymes within the hydrogel (Traffano-Schiffo et al., 2017a).

The alginate microstructure depends on several factors such as the alginate's concentration, average molecular weight and monomer composition (M/G ratio), as well as the presence of secondary excipients and the synthesis conditions (mainly the  $\text{Ca}^{2+}$  concentration and pH). On the other hand, the alginate bead constitutes an intrinsically inhomogeneous system, determined by the dropping method followed in conventional synthesis procedures. This method generates a  $\text{Ca}^{2+}$  gradient established from the surface to the core of the forming bead (Thu et al., 2000). All these parameters are related to the capacity of the hydrogel network to interact with the encapsulated biomolecule and the surrounding medium (Santagapita et al., 2011; Gombotz & Wee, 2012).

One of the most precise and powerful techniques to evaluate the microstructure of hydrogels is the small-angle X-ray scattering (SAXS). The SAXS method is able to reveal subtle differences in electron density within hydrogels cross-linked networks in the range 1-100 nm, providing information on the supramolecular structure formed by biopolymers (Waters et al., 2010). SAXS patterns are indicative of rod like objects formed as the accumulation of cross-linked alginate chain dimers randomly orientated, establishing junction zones of different multiplicity (Agulhon, Robitzer, David, & Quignard, 2012). Previous works of lactase encapsulation in Ca(II)-alginate beads (Traffano-Schiffo et al., 2017a; Traffano-Schiffo, Aguirre Calvo, Castro-Giraldez, Fito, & Santagapita, 2017b) revealed that the remaining lactase activity after storage, freezing and freeze/thaw cycles, and dehydration, preserved by trehalose, was even improved by the presence of gums. The microstructure of the

beads generated at pH 3.8 showed rods with smaller cross-sectional radius and with lower compactness when gums were used as additives. However, there is no available information on how the microstructure is affected due to the constraints imposed by the operation conditions. Thus, the aim of this research is to study the microstructural changes of Ca(II)-alginate matrices driven by the presence of trehalose, arabic and guar gums as excipients and to correlate these changes with the enzymatic activity of the encapsulated lactase in operation conditions.

## **2. Materials and Methods**

### **2.1. Materials**

The employed materials are listed below: sodium alginate (Algogel 5540) from Cargill S.A. (San Isidro, Buenos Aires, Argentina), molecular weight of  $1.97 \cdot 10^5$  g/mol and mannuronate/guluronate ratio of 0.6; D-trehalosedihydrate (Hayashibara Co., Ltd., Shimoishii, Okayama, Japan/Cargill Inc., Minneapolis, Minnesota, USA), molecular weight of 378 g/mol; guar gum (Cordis S.A., Villa Luzuriaga, Buenos Aires, Argentina), molecular weight of 220.000 g/mol and a mannose/galactose ratio of 1.8; arabic gum (Biopack, Zárate, Buenos Aires, Argentina), molecular weight of 250.000 g/mol and a purity of 99%;  $\beta$ -galactosidase (lactase) from *Aspergillus Oryzae* (8.0 U/mg) (Sigma-Aldrich Co., Ltd., Saint Louis, USA). One enzymatic unit was defined as the amount of enzyme able to hydrolyze 1.0  $\mu$ mol of lactose per minute at pH 4.5 at 30°C.

### **2.2. Gel Beads Preparation**

Four different formulations for the encapsulation of the enzyme (E) were prepared, with the following composition: alginate (EA); alginate-trehalose (EAT); alginate-trehalose-guar gum (EATGG); alginate-trehalose-arabic gum (EATAG). All the solutions were prepared in 0.1 M acetate buffer pH 3.8. The final concentration of lactase was 0.775 mg/mL. The enzyme and the precursor solutions were carefully mixed and maintained at  $4 \pm 1$  °C in order to avoid enzyme activity losses. Taking into account that the isoelectric point of the enzyme is 4.61 (Dashevsky, 1998) and the  $pK_a$  values of alginate are 3.38 and 3.65 (Smidsrød, Larsen, Painter, & Haug, 1969), the buffer acetate at pH 3.8 was used in order to generate favorable electrostatic interactions between the negatively charged alginate and the positively charged enzyme. A peristaltic pump was used to drop 10 mL of the alginate-enzyme mixture into 100 mL of the gelling solution, according to the drop method described by Austin, Bower and Muldoon with some modifications (Austin, Bower, & Muldoon, 1996). For EA beads preparation, 1% (w/v) alginate solution containing lactase was dropped into 2.5% (w/v)  $\text{CaCl}_2$  solution prepared in 0.1 M acetate buffer pH 3.8. For EAT, EATGG, and EATAG preparation, a 1% (w/v) alginate with 20% (w/v) trehalose with or without 0.25% (w/v) of guar or arabic gums containing the enzyme was dropped into the 2.5% (w/v)  $\text{CaCl}_2$  solution supplemented with 20% (w/v) trehalose using the same procedure previously described.

The  $\text{CaCl}_2$  solution (with or without trehalose) was maintained in a cold bath with constant stirring. A needle with 0.25 mm diameter and 6 mm length (Novofine 32 G, Novo Nordisk A/S, Bagsvaerd, Denmark) was used for the dropping. The pump speed was  $9.0 \pm 0$  rpm. The distance between the needle and the  $\text{CaCl}_2$  solution was 6.0 cm. After beads generation, they were maintained for 15 min in  $\text{CaCl}_2$  solution

(with constant stirring), and then they were washed 5 times with bidistilled cold water ( $5\pm 1$  °C) in order to remove free  $\text{Ca}^{2+}$ . Previous to SAXS measurements, beads were maintained at ( $5\pm 1$  °C) for 96 h.

Control beads without enzyme were also prepared using the same methodology.

### 2.3. Microstructure Analyses

Beads synthesized at pH 3.8 were incubated in 0.1 M acetate buffer pH 4.5 for different times (5, 10, 15, 20, 40, and 60 min). The microstructure characterization was performed by SAXS at the LNLS SAXS2 beamline in Campinas, Brazil, working at  $\lambda = 0.1488$  nm. The wave vector range was selected in the range  $0.096 \text{ nm}^{-1} < q < 2.856 \text{ nm}^{-1}$ . All the Ca(II)-alginate beads analyzed showed isotropic scattering and were modeled as a fractal system composed of rodlike structures, although the rigorous interpretation of experimental results as indicating “fractality” requires many orders of magnitude of power-law scaling (Sonego et al., 2016). Five parameters were analyzed: i)  $\alpha_1$ , the fractal dimension at distances higher than the characteristic size of the rods composing the structure ( $R_1$ ), which describes the multiplicity of the junction zone, at  $q < 0.28$ ; ii)  $\alpha_2$ , the fractal dimension at distances lower than  $R_1$ , at  $q > 0.55$ , describing the degree of compactness within the rods; iii)  $R_1$ , the outer radius of the fibrils, which is given by  $R_1 = R_g\sqrt{2}$ ,  $R_g$  being the mean gyration radius in the cross-section of the rods, which is obtained from the maximum exhibit by the Kratky plot at  $q \approx 1/R_g$ ; iv)  $\alpha_3$ , related to the connectivity between associated polymer chains forming dimers (basic units composing the rods) at  $q > 1.5$  and v)  $R_2$ , related to the outer radius of these aforementioned units.



The Kratky plot: scattering intensity multiplied by the square modulus of the scattering vector,  $I(q) \cdot q^2$ , as a function of the modulus of the scattering vector,  $q$ , gives a maximum value at the intersection of the  $\alpha_1$  and  $\alpha_2$  power law regions and a minimum value at the intersection of the  $\alpha_2$  and  $\alpha_3$  power law regions, allowing the calculation of parameters  $R_1$  and  $R_2$ , respectively. All measurements were made in triplicate. Parameters  $\alpha_1$ ,  $\alpha_2$  and  $\alpha_3$  were evaluated from the slope of the scattering intensity averaged along azimuthal angles versus the scattering vector  $q$  in the log–log scale.

#### 2.4. $\beta$ -Galactosidase Activity

The enzyme activity was evaluated based on the absorbance values at 420 nm by using a Jasco V-630 UV-vis spectrophotometer (JASCO Inc., Maryland, USA) at room temperature and following the method described by Park, Santi, and Pastore, with some modifications (Park, Santi, & Pastore, 1979).

Each analysis was performed using a pool of 9 beads which were mixed with 0.25 mL of 0.25% (w/v) *o*-nitrophenyl- $\beta$ -D-galactopyranoside (ONPG) (Sigma Chemical Co.) prepared with 0.1 M acetate buffer pH 4.5 and incubated during 5, 10, 15, 20, 40, and 60 min at 33 °C, without stirring in order to avoid possible enzymatic activity losses. The reaction was stopped adding 0.5 mL of sodium carbonate 10% (w/v) and 0.25 mL of sodium citrate 10% (w/v) was added to dissolve the beads and finally, 1.75 mL of distilled water was added for subsequent measurement of *o*-nitrophenol (ONP) at 420 nm. Measurements were made in triplicate.

The effect of the composition of the beads on the enzyme stability was evaluated through an activity index calculated with the following equation:

$$Activity\ Index\ (dimensionless) = \frac{Activity_t}{Activity_0} \quad (1)$$

where  $Activity_t$  corresponds to the activity of any of the studied systems (EA, EAT, EATAG or EATGG) at time  $t$  (5, 10, 15, 20, 40, or 60 min) and  $Activity_0$  is the activity of the same system determined as described by Traffano-Schiffo et al. (2017b).

### 2.5. Low Field Nuclear Magnetic Resonance (LF-NMR)

Diffusion coefficient of water ( $D_w$ ) was measured by time resolved low field proton nuclear magnetic resonance ( $^1H$ -LF-NMR) in a Bruker Minispec mq20 (Bruker Biospin GmbH, Rheinstetten, Germany) with a 0.47 T magnetic field operating at a resonance frequency of 20 MHz. Two samples of each bead system were previously equilibrated at  $25.00 \pm 0.01$  °C in a thermal bath (Haake, model Phoenix II C35P, Thermo Electron Corporation GmbH, Karlsruhe, Germany).

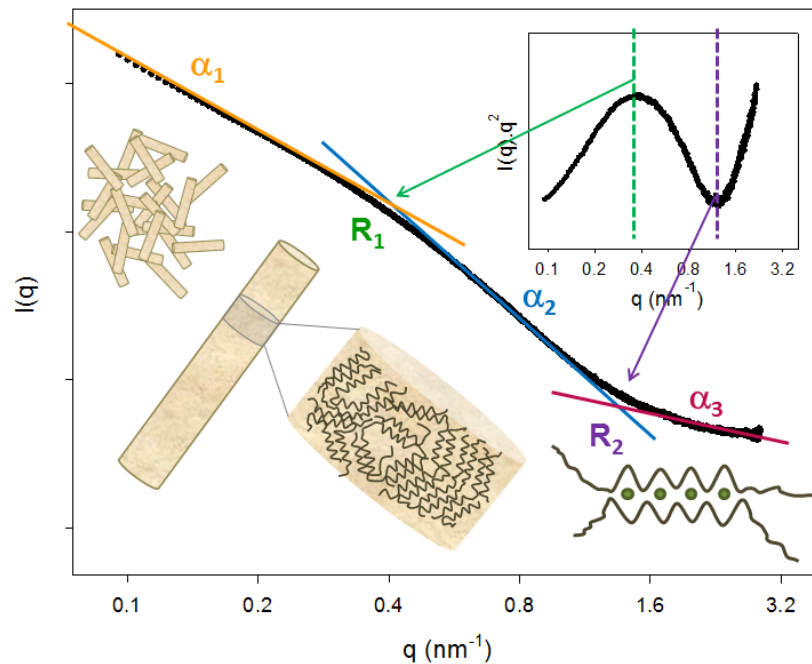
Measurements were performed through the pulsed magnetic field gradient-spin echo sequence (Stejskal, & Tanner, 1965). The applied magnetic field gradient intensity was calibrated in the range between 1.4 and 2.4 T/m by employing 1.25 g/L  $CuSO_4 \cdot 5H_2O$  water solution, characterized by a known  $D_w$  value ( $2.3 \cdot 10^{-9}$  m<sup>2</sup>/s at 25 °C) (Hester-Reilly, & Shapley, 2007). The bead samples were analyzed by setting the magnetic field gradient amplitude to 1.4 T/m,  $t$  (the time between 90 and 180 pulses) to 7.5 ms,  $\delta$  to 0.5 ms, and  $\Delta$  to 7.5 ms. The number of scans, the recycle delay and gain were 16, 2 s and 69 dB, respectively.  $D_w$  was calculated following the procedure reported by Santagapita et al. (2013).

### 2.6. Statistical Analyses

The statistical analyses were performed by one-way ANOVA with Tukey's post test by using Prism 6 (GraphPad Software Inc., San Diego, CA, USA) in order to determine significant differences between the mean values of beads of different compositions on the measured parameters.

### **3. Results and Discussion**

Figure 1 shows the scattering intensity as a function of the scattering vector for a representative sample of Ca(II)-alginate hydrogel containing lactase (EA), incubated in 0.1 M acetate buffer pH 4.5 for 20 minutes. A schematic representation of the scale of different structural parameters derived from SAXS scattering experiments is included. SAXS scattering curves can be divided in three regions: at low, intermediate and high  $q$  values. From the slope of the log-log plot at low  $q$ ,  $\alpha_1$  parameter is indicative of the interconnectivity of the rods composing the alginate network. At intermediate  $q$  values,  $\alpha_2$  parameter represents the compactness within the rods, and at high  $q$  values  $\alpha_3$  parameter characterizes the connectivity between associated polymer chains forming dimers. Besides, two characteristic radii of gyration can be derived:  $R_1$ , related to the multiplicity of the junction zone domains in the alginate rod, and  $R_2$  denoting the size of the polymer dimers basic units. Kratky plots ( $q^2 \cdot I(q)$  vs  $q$ ) were used to evaluate  $R_1$  as already reported (Sonego et al., 2016; Thu et al., 2000; Gombotz & Wee, 2012; Waters et al., 2010; Agulhon et al., 2012). Characteristic SAXS profiles and Kratky plots from all the systems analyzed are presented in Supplementary File (Figures S1 and S2, respectively).



**Figure 1.** log–log SAXS profile plot of a representative sample of Ca(II)-alginate hydrogel containing lactase (EA), incubated in 0.1 M acetate buffer pH 4.5 for 20 minutes. Parameters  $\alpha_1$ ,  $\alpha_2$  and  $\alpha_3$  were evaluated from the slope of the scattering intensity at low, intermediate and high  $q$  values, respectively. The radius of gyration of the rods (parameter  $R_1$ ) and the characteristic size of the Ca(II)-alginate dimers (parameter  $R_2$ ) was obtained from the Kratky plot (inset).

Control systems without the enzyme were also conducted in order to analyze the influence of the presence of lactase in the microstructure of the alginate gel. Except for subtle changes, the microstructure remains the same for all the systems under study and thus it can be concluded that the enzyme is not directly involved in the formation of the alginate rods. A complete description of these results is included in Supplementary File (Figure S3).

**Figure 2.** Microstructure parameters of Ca(II)-alginate beads synthesized at pH 3.8 vs. time of incubation in 0.1 M acetate buffer pH 4.5. a. Rod cross-sectional radius ( $R_1$ ) deduced from the maxima obtained on Kratky plots. b. Fractal dimension at distances lower than  $R_1$  or parameter  $\alpha_2$  of the microstructure derived from log-log SAXS profiles. Standard deviations values are included. E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum.

Figure 2 shows the evolution of the microstructure parameters related to the size and compactness of the rods ( $R_1$  and  $\alpha_2$ , respectively). For the non-incubated beads (synthesized at pH 3.8), the addition of trehalose with or without gums affected the extent of the rod formation, reducing the outer radius of the fibrils (Figure 2a, time = 0) as well as their compactness (Figure 2b, time = 0), as previously discussed Traffano-Schiffo et al. (2017b).

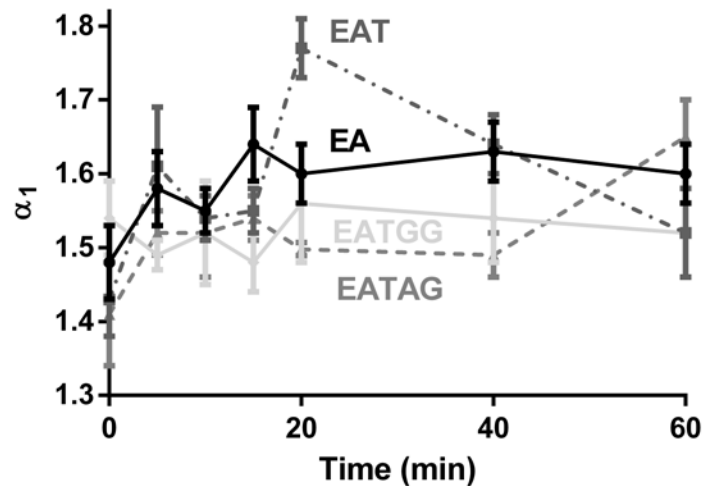
As a general trend, the increase in pH (from 3.8 to 4.5) generates a decrease both in size and compactness of the rods for the systems in the absence of gums. However, the presence of guar or arabic gums generates a much stable structure, maintaining a constant rod size and significantly reducing the initial drop in  $\alpha_2$ , as well as further changes along incubation time. It is worth noting that the synthesis pH is close to the isoelectric point of alginate and thus the contribution of H-alginate to the

consolidation of the alginate network cannot be neglected. Therefore, at low free  $\text{Ca}^{2+}$  concentrations, an increase in pH would lead to an increase in alginate polymer repulsion with a concomitant reduction of the compactness due to the loss of the H-alginate contribution to the structure. This can also explain the reduction in the rod size, considering that some of the alginate chains can be released from the junction zones. The presence of gums seems to reduce alginate-alginate repulsion which suggests that they are located in within the rods, particularly between associated polymer chains forming dimers. This hypothesis was deeply analyzed by considering the evolution of  $\alpha_3$  and  $R_2$ , the nanostructure parameters describing the size and compactness of these dimers, basic units of construction of the rods (as shown in Figure 3 a and b, respectively). The presence of gums modulates the nanostructure of the dimers generating an increase in their size and a decrease in their compactness, compatible with the hypothesis of gums being intercalated within the dimers. Furthermore, the addition of gums once again induced a higher stability toward an increase in pH. However and regardless of the presence of different excipients, the nanostructure of dimers shows similar characteristics and remains stable during the operation conditions at pH 4.5.

**Figure 3.** Microstructure parameters of Ca(II)-alginate beads synthesized at pH 3.8 vs. time of incubation in 0.1 M acetate buffer pH 4.5. a. Characteristic size of the Ca(II)-alginate dimers ( $R_2$ ) deduced from the minima obtained on Kratky plots. b. Fractal dimension at distances lower than  $R_2$  or parameter  $\alpha_3$  of the microstructure derived from log-log SAXS profiles. Standard deviations values are included. E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum.

In order to go a step further in the analysis of the microstructure of the hydrogels during the operation conditions, the evolution of  $\alpha_1$  parameter for each bead composition was analyzed (Figure 4). The parameter  $\alpha_1$  is related with the degree of interconnection of the rods composing the structure. Values of  $\alpha_1$  around 1 correspond to randomly oriented rods, while a  $\alpha_1$  close to 2 is indicative of a network composed of well interconnected (Sonego et al., 2016). All the systems under study showed initial values of  $\alpha_1$  around 1.5 (i. e. for the hydrogel obtained at pH 3.8), characteristic of a consolidated network. The formulations lacking gums additives showed an increase in  $\alpha_1$  when exposed to the operation pH 4.5, reaching values of around 1.8 (for the system with trehalose, EAT, after 20 minutes of incubation). Once established the degree of interconnectivity of rods in aged systems, the  $\alpha_1$

parameter is expected to remain constant in time. One plausible explanation for  $\alpha_1$  fluctuations relies on the increased availability of alginate polymer chains resulting from the partial dissolution of rods, as previously discussed regarding parameter  $\alpha_3$  and  $R_2$ . Thus, the presence of free alginate chains could induce the formation of new connections in the network establishing a new equilibrium. In line with this, the presence of gums induces stabilization in the microstructure, resulting in no significant variations in  $\alpha_1$  parameter both with the initial increase in pH and throughout the operation time at pH 4.5.



**Figure 4.** Fractal dimension at distances higher than  $R$  or parameter  $\alpha_1$  of the microstructure derived from log-log SAXS profiles. Standard deviation values are included. E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum.

Water self-diffusion coefficient ( $D_w$ ) determined by LF-NMR for Ca(II)-alginate beads gives valuable information of the overall decrease on the mobility of protons (especially those from water), modulated by the exchange between them and the protons of the biopolymers (Sonego et al., 2016; Aguirre Calvo, Busch, &



Santagapita, 2017). The magnitude of this decrease depends on reduced flexibility of the biopolymer chains with respect to water and the aggregation state. As expected, all alginate systems gelled at pH 3.8 showed reduced  $D_w$  values in comparison with water,  $(2.30 \pm 0.01) \times 10^{-9} \text{ m}^2 \text{ s}^{-1}$  (Holz, Heil, & Sacco, 2000), as shown in Table 1. The presence of trehalose in the hydrogels further reduced  $D_w$  values. However, the concomitant addition of gums induced a slight increase of  $D_w$  with respect to the EAT system. These changes are in well agreement with the water activity and content previously reported (Traffano-Schiffo et al., 2017b). Besides, the measure of  $D_w$  can give an idea of the water transport through the aqueous macropores, which can be also related with the diffusion of other water soluble species. The dramatic drop in  $D_w$  can be explained from the evolution of parameters  $\alpha_2$  and  $R_1$ , which show a significantly higher rod size and compactness for the hydrogel in the absence of additives (EA). This is consistent with highly packed rods which in turn enclose more alginate dimers, reducing the tortuosity through the aqueous macropores.

**Table 1.** Diffusion coefficients of water ( $D_w$ ) in beads containing lactase determined by LF-NMR at 25 °C.

<b>System</b>	<b><math>D_w</math> (<math>10^{-9} \text{ m}^2/\text{s}</math>)</b>
<b>EA</b>	2.1 ± 0.03 <sup>a</sup>
<b>EAT</b>	1.492 ± 0.011 <sup>c</sup>
<b>EATAG</b>	1.6 ± 0.02 <sup>b</sup>
<b>EATGG</b>	1.613 ± 0.013 <sup>b</sup>

<sup>a-c</sup> E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum. Standard deviation values are included. Different letters on the columns indicate significant differences between means ( $p < 0.05$ ).

Then, bearing in mind that the modulation of the enzyme activity will be given by the entrance of substrate through the macropores as well as by the overall microstructure of the Ca(II)-alginate gel, the addition of excipients would be expected to modulate lactase activity. Figure 5 shows the effect of the composition of the beads on  $\beta$ -galactosidase stability by measuring the ONP concentration (colored compound) spectrophotometrically. It can be observed that the EAT shows the lowest ratio of enzymatic activity; however the addition of gums as second excipients reverts this detriment caused by trehalose, increasing for guar gum containing beads (EATGG)  $\beta$ -galactosidase activity above the standards of bear Ca(II)-alginate hydrogel (EA). EAT beads present the worst possible scenario: low diffusion and poor microstructure stability. The incorporation of gums palliates this by improving the intrinsic transport properties of the initial hydrogel, in one hand, and more importantly by stabilizing its microstructure. The latter seems to be a determining factor in the performance of these systems during operating conditions.

#### **4. Conclusions**

Though being a key in preserving enzymatic activity toward freezing and dehydration processes, the addition of trehalose as additive in Ca(II)-alginate lactase encapsulation systems prompts important structural changes leading to a loss of enzymatic activity during operation conditions. Trehalose drastically reduces the water self-diffusion coefficient in the starting hydrogel system (synthesized at pH 3.8), concomitant with the reduction in size and compactness of the structure at the scale of alginate rods. The incorporation of gums as second excipients does not

revert this last effect, nevertheless induces a significant stabilization in the microstructure not only at the rod scale, but also at smaller and bigger scales. On one hand, gums are shown to induce an important modulation in the characteristic size and density of alginate dimers (basic units of construction of rods). On the other hand and highly related to this, the variation in the degree of interconnection of rods evidenced in EA and EAT systems (i.e. in the absence of gums) is impeded. It is worth noting that in EA and EAT, this process is facilitated by free alginate polymer chains that become available from partial rod dissolution. Thus, this microstructure stabilization induced by the concomitant addition of guar gum results in an increased lactase activity during operation conditions, improving the performance of the systems with trehalose even above the bear Ca(II)-alginate matrix.

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### **Appendix A. Supplementary data**

Figure S1. log–log SAXS profile plots of representative of Ca(II)-alginate lactase beads for the different systems, incubated in 0.1 M acetate buffer pH 4.5 at different

times. E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum. Figure S2. Kratky plots of SAXS data obtained for Ca(II)-alginate beads containing enzyme for the four systems analyzed. The curves at each incubation time were arbitrary shifted up. E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum. Figure S3. Parameter  $\alpha_1$  (a) and  $\alpha_2$  (b) of the microstructure derived from log-log SAXS profiles and  $R_1$  (outer radius of the rods) (c), deduced from the maxima obtained on Kratky plots. Standard deviations values are included. Different letters on the columns (a-e) indicate significant differences between values with  $p < 0.05$ . E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum.

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### **Highlights**

- Trehalose plays a key role in preserving enzymatic activity against thermal treatments
- Trehalose reduces the water self-diffusion coefficient in Ca(II)-alginate beads
- Gums as second excipients induce a stabilization in the microstructure at rod scale
- Gums stabilize the size and density of the dimers and the interconnection of rods
- Gums improve the lactase activity preservation

**GUMS INDUCED MICROSTRUCTURE STABILITY IN Ca(II)-ALGINATE  
BEADS CONTAINING LACTASE ANALYSED BY SAXS**

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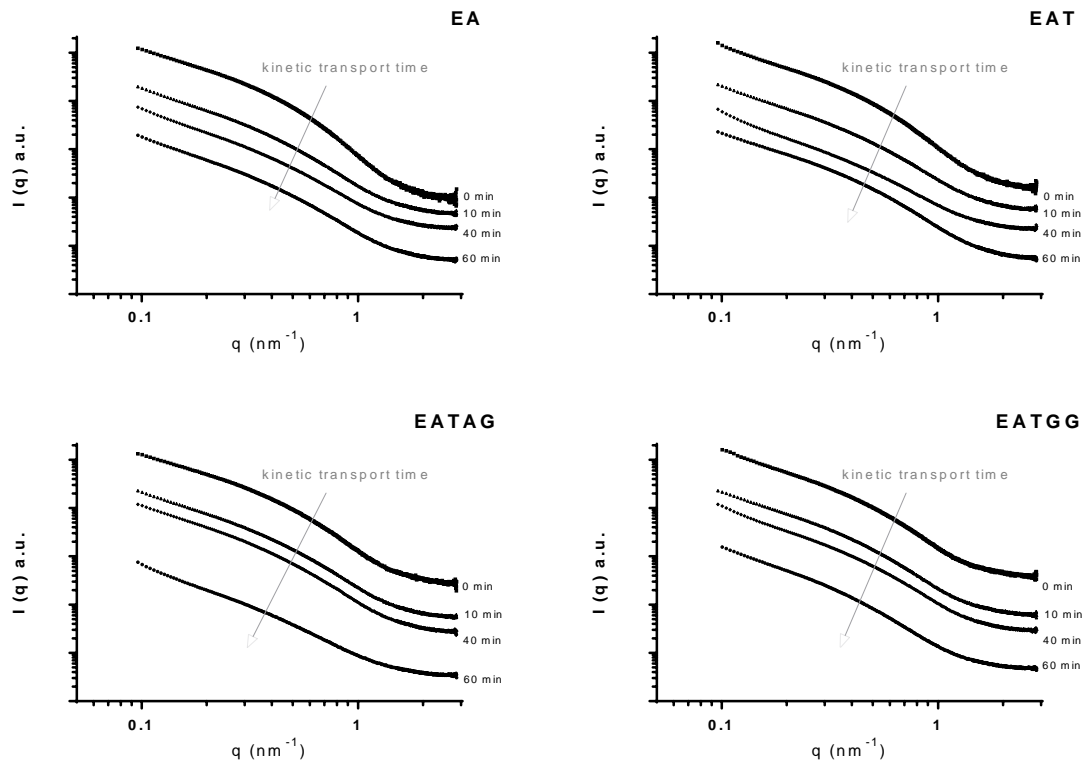
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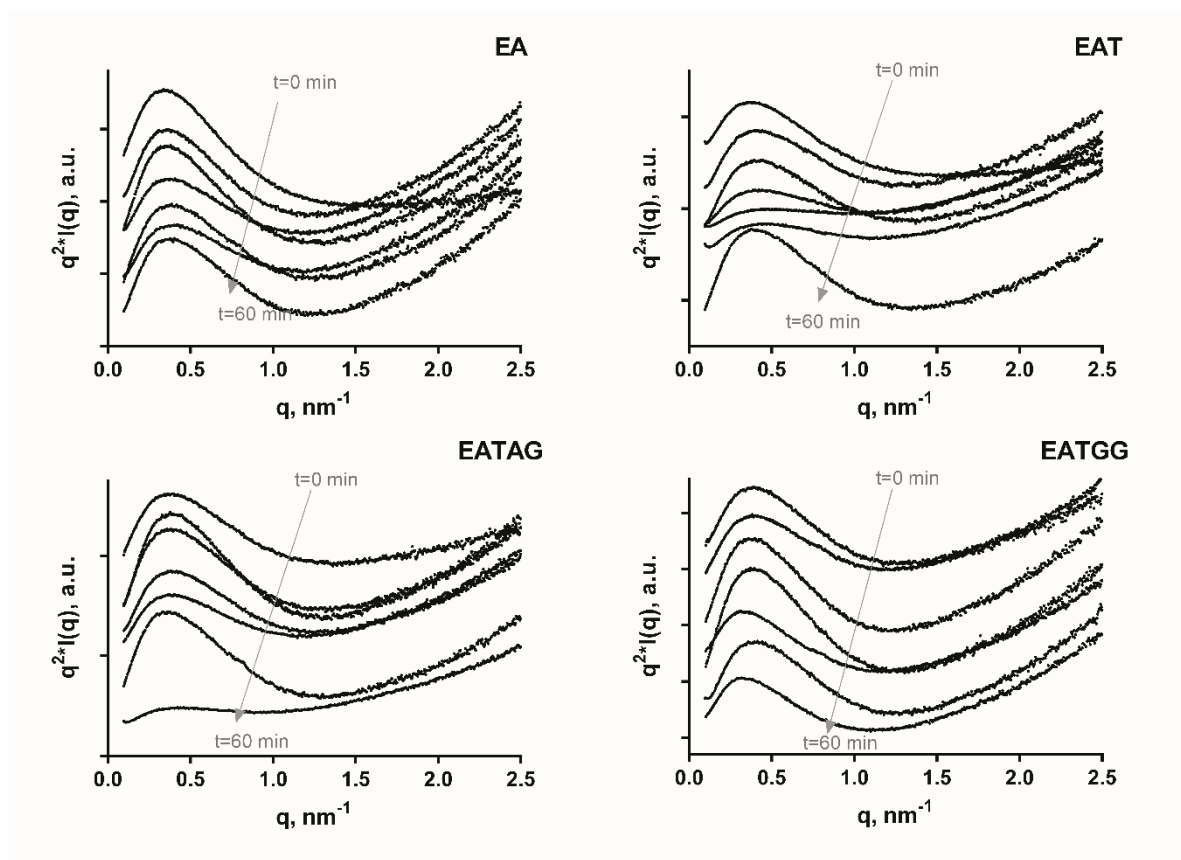
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SUPPLEMENTARY FILE

Characteristics SAXS profiles and Kratky plots at different times of operation conditions from all the systems analyzed are presented in Figures S1 and S2, respectively).



**Figure S1.** log–log SAXS profile plots of representative of Ca(II)-alginate lactase beads for the different systems, incubated in 0.1 M acetate buffer pH 4.5 at different times. E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum.

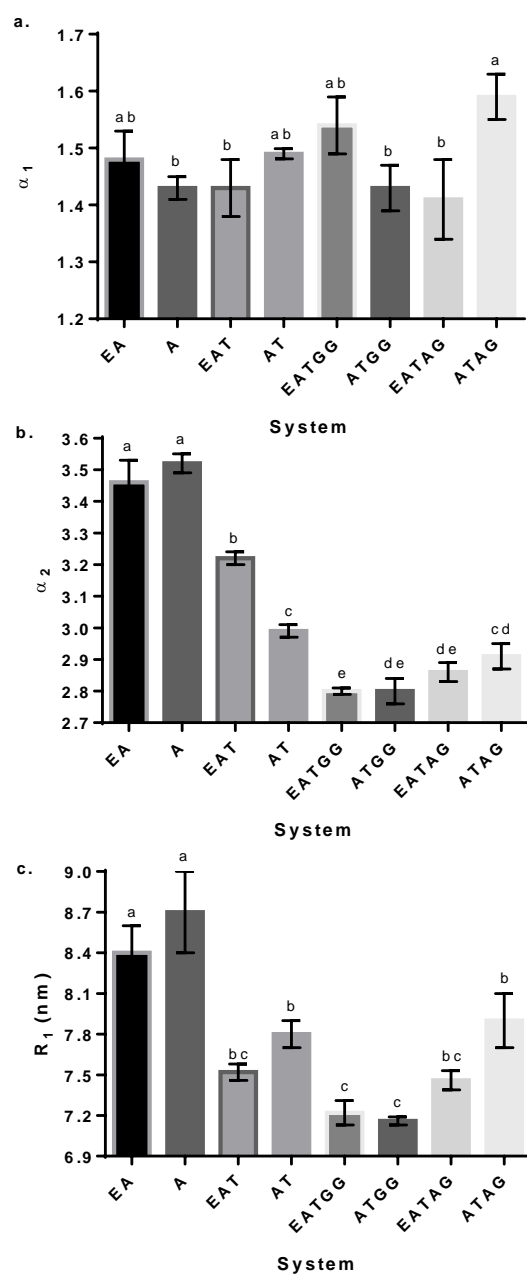


**Figure S2.** Kratky plots of SAXS data obtained for Ca(II)-alginate beads containing enzyme for the four systems analyzed. The curves at each incubation time were arbitrary shifted up. E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum.

Figure S3 shows the  $\alpha_1$ ,  $\alpha_2$  and  $R_1$  parameters obtained from the control samples prepared without lactase. Minor changes were observed between the different systems with and without lactase. For instance: the interconnectivity of the rods in the presence of arabic gum was higher in the absence of enzyme. However, neither the rods size nor its compactness was affected by the enzyme in this system. Also, the compactness of the rods in the presence of trehalose was higher in the presence of the enzyme, but the rods size remains unmodified (and showed even lower mean

values). If the enzyme is directly involved in the formation of the alginate rods, a change in both size and compactness of the rods is expected, which in turn could affect the interconnection between rods at a larger scale. Then, since slightly changes were observed, we conclude that the microstructure remains the same for all the systems under study in presence or absence of the enzyme.

A complete analysis of the observed changes between the beads containing the different additives can be read in Traffano-Schiffo et al., 2017.



**Figure S3.** Parameter  $\alpha_1$  (a) and  $\alpha_2$  (b) of the microstructure derived from log-log SAXS profiles and  $R_1$  (outer radius of the rods) (c), deduced from the maxima obtained on Kratky plots. Standard deviations values are included. Different letters on the columns (a-e) indicate significant differences between values with  $p < 0.05$ . E, enzyme; A, alginate; T, trehalose; AG, arabic gum; GG, guar gum.

