

Insights on the Interactions of Chitosan with Phospholipid Vesicles. ² Part I: Effect of Polymer Deprotonation

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S Supporting Information 6

ABSTRACT: Interactions between the polysaccharide chitosan and negatively charged 7 8 phospholipid liposomes were studied as a function of compositional and environmental conditions. Using isothermal titration calorimetry, different levels of deprotonation of 9 chitosan in acidic solutions were attained with titration of the fully protonated polymer at 10 pH 4.48 into solutions with increasing pH. The process was found to be highly 11endothermic. We then examined the interaction of the polymer with vesicles in solutions 12 of different pH. Even when partially deprotonated, the chitosan chains retain their affinity 13



to the negatively charged liposomes. However, the stronger adsorption results in lower organization of the chains over the 14 membrane. 15

1. INTRODUCTION

16 Lipid vesicles represent a variety of colloidal structures formed 17 by the self-assembly of phospholipid molecules into bilayers. 18 Liposomes with sizes in the 100 nm range are probably the 19 most studied and well developed example of nanovesicles 20 applied as advanced drug delivery systems.¹⁻¹⁰

Chitosan is a well-known polysaccharide, obtained from 21 22 deacetylation of natural chitin which is the main component of 23 crustacean shells. Currently, chitosan is being considered as an 24 important biocompatible macromolecule¹¹ extensively used for 25 the development of drug and vaccine carriers aiming to 26 optimize efficiency of treatments and achieve controlled antigen $_{27}$ release. $^{12-20}$ The polymer is soluble in aqueous solutions with 28 pH lower than 5.5 as a result of protonation of the amino 29 groups along the polymer chain; see Figure 1A. The 30 protonation results in an extended polyelectrolyte which can 31 exhibit strong interactions with phospholipids in model and 32 cellular membranes²¹⁻²⁶ as well as with negatively charged 33 surfaces.²⁷ Hence, the mucoadhesive property of chitosan has 34 been explored in the development of chitosomes (liposomes 35 modified with chitosan) with the aim not only to target drug 36 carriers to specific sites of action, but also to further prolong the 37 drug therapeutic action by keeping the vesicles adsorbed over 38 the cell membranes for increased periods of time.^{28,29}

Nevertheless, the well-known instability of drug delivery 39 40 systems in biological media prevents the efficiency of the 41 carriers. Thus, the development of such systems requires 42 detailed knowledge of all physical, chemical, and biological 43 characteristics playing a role in processes involved at many 44 levels, starting from the production to the final in vivo 45 performance. In what concerns liposomes modified with 46 chitosan, in the recent years, deeper understanding has been 47 gained on fundamental aspects related to the vesicle ⁴⁸ preparation method,³⁰ physical and morphological features,^{31,32} 49 stability parameters under temperature variation,³³ and

molecular interactions.³⁴ The applicability of these vesicles as 50 a vaccine delivery system has also been evaluated.^{12,14} More 51 recently, we have studied the thermodynamic characteristics of 52 the binding of chitosan onto the membranes of liposomes made 53 of zwitterionic phospholipids and different fractions of 54 negatively charged phospholipids.³⁵ The electrostatic inter- 55 actions between the macromolecules and the membrane were 56 found to be regulated by and to increase with the vesicles 57 surface charge. At the point of saturation via charge 58 compensation, the vesicles were shown to aggregate. The role 59 of polymer charge, that is, degree of deionization or 60 deprotonation, was not investigated.

To further elucidate the processes occurring during the 62 encounter of chitosan with phospholipids vesicles, our purpose 63 in the present report was to explore the effect of deprotonation 64 of the polymer amino groups as a driving force governing the 65 interaction. We first studied in more detail and thermodynami- 66 cally characterized the deprotonation of chitosan introduced in 67 solutions of various pH. We then performed a thorough 68 investigation of the effect of deprotonation on the polymer 69 stability and interaction with partially charged lipid vesicles. 70 Having characterized the properties of the system from the 71 viewpoint of the polysaccharide, in a following study, we 72 pursued the characterization of the membrane response in 73 terms of mechanical properties and stability upon contact with 74 chitosan.36 75

2. MATERIALS AND METHODS

2.1. Materials. Chloroform solutions of 1,2-dioleoyl-sn-glycero-3- 76 phosphatidylcholine (DOPC) and 1,2-dioleoyl-sn-glycero-3-phospha-77 tidylglycerol (sodium salt) (DOPG) were purchased from Avanti Polar 78

Received: August 20, 2013 **Revised:** October 28, 2013 79 Lipids Inc. (Birmingham, AL) and used without further purification. 80 They were stored at -20 °C upon arrival. Chitosan was a gift from 81 Primex (Germany), with 95% degree of deacetylation (DDA). The 82 average molecular weight was determined as $M_w = 199$ kDa 83 (corresponding to 1223 repeat monomers per molecule) by multiangle 84 laser light scattering size exclusion chromatography (MALLS-SEC),³⁷ 85 with a radius of gyration of 46 nm.

86 All other reagents were of analytical grade. All solutions were 87 prepared using deionized water from Milli-Q Millipore system with a 88 total organic carbon value of less than 15 ppb and a resistivity of 18 89 M Ω cm.

2.2. Preparation of Large Unilamellar Vesicles (LUVs) and 90 91 Chitosan Solutions. The lipid solutions in chloroform were 92 transferred into round-bottom flasks, and the organic solvent was 93 removed by evaporation under a stream of nitrogen gas until complete 94 drying followed by 2 h in a vacuum desiccator. An aqueous buffer 95 solution of acetic acid/sodium acetate was added in the flask with the 96 lipid film. All buffers were prepared at a total concentration of 80 mM, 97 but varying weight fractions of acid and salt in order to obtain the 98 desired pH: 4.90, 5.30, 6.00, or 7.10 with a maximal variation of 0.01. 99 Liposomes were obtained by vortexing for about 2 min, followed by 100 extrusion using a LipsosoFast pneumatic extruder (Avestin Inc., 101 Ottawa, Canada) operating at a pressure of 200 kPa. The final total 102 lipid concentration in all experiments was 3.82 mM and the molar 103 ratio between DOPC and DOPG was adjusted to obtain negatively 104 charged vesicles with 10 mol % DOPG (0.38 mM). The extrusion was performed in three consecutive steps: 20 times extrusion through a 105 106 400 nm diameter pore polycarbonate filter, 20 times through a 200 nm 107 diameter pore filter, and finally 40 times through a 100 nm diameter pore filter. Vesicles prepared in this way generally have a narrow size 108 109 distribution as confirmed with dynamic light scattering (PDI around 110 1.5) and are known to be almost entirely unilamellar.

The chitosan solution was prepared by vigorous overnight stirring 112 of the powder in the acetate buffer (pH 4.48 \pm 0.01), at a 113 concentration of 1 mg/mL. Solutions with lower concentrations were 114 prepared by diluting the stock solution with buffer. The pH of all 115 solutions was constantly monitored before and after sample 116 preparation and the conductivity was measured, to ensure constant 117 ionic strength. Acetic acid was carefully added to adjust the pH when 118 required.

2.3. Dynamic Light Scattering and *ζ***-Potential.** Size distribution measurements on chitosan solutions and LUV suspensions were performed with Zetasizer Nano ZS instrument (Malvern Instruments, Worcestershire, U.K.). The instrument uses a 4 mW HeNe laser at a wavelength of 632.8 nm and detection at an angle of 173°. All measurements were performed in a temperature controlled chamber at 25 °C. The autocorrelation function was acquired using exponential spacing of the correlation time. The data analyses were performed with provided by Malvern. The intensity-weighted size distribution was obtained by fitting data with a discrete Laplace inversion routine.³⁹

129 The ζ -potential of vesicles and chitosan was analyzed in the same 130 Malvern Zetasizer instrument performing at least six runs per sample. 131 The measurement principle is based on laser Doppler velocimetry. 132 The electrophoretic mobility u is converted to ζ -potential using the 133 Helmholtz–Smoluchowski relation $\zeta = u\eta/\varepsilon\varepsilon_0$, where η is the solution 134 viscosity, ε the dielectric constant of water, and ε_0 the permittivity of 135 free space.

2.4. Isothermal Titration Calorimetry (ITC). ITC measurements 136 137 were performed with a VP-ITC microcalorimeter from MicroCal Inc. (Northampton, MA). The working cell (1.442 mL in volume) was 138 139 filled with the LUV suspension, and the reference cell with the 140 corresponding liposome-free buffer solution. One aliquot of 2 μ L 141 followed by 27 aliquots of 10 μ L of chitosan solution (pH 4.48 ± 0.01) 142 were injected stepwise with 200 s intervals into the working cell filled 143 with the vesicle suspension of variable pH, namely, 4.90, 5.30, 6.00, 144 and 7.10 (± 0.01). The corresponding reference experiments were also 145 performed, that is, titration of chitosan-free buffer in vesicle suspension 146 and titration of chitosan solution in vesicle-free buffer. To avoid the 147 presence of bubbles, all samples were degassed for 10 min shortly 148 before performing the measurements. The sample cell was constantly stirred at a rate of 307 rpm, and the measurements were performed at 149 25 °C. The data analyses were carried out with Origin software 150 provided by MicroCal. 151

3. RESULTS AND DISCUSSION

3.1. Chitosan Deprotonation. Intuitively, electrostatic 152 interactions are first to consider when discussing binding of the 153 positively charged chitosan to negatively charged vesicles. 154 Considering the dimensions of the (small) vesicles and the long 155 chitosan backbone, the polymer has to bend when adsorbing to 156 the LUV membrane, altering the surface charge of the vesicles. 157 Apart from these effects, we considered another factor 158 promoting the interaction between the two identities, namely, 159 the deprotonation of the chitosan chains or, in other words, the 160 transfer of protons from some amino groups to the solution. 161 Deprotonation was found to play a role in the binding of 162 chitosan to DNA.40 Associated effects may be insignificant in 163 conditions where chitosan is permanently protonated, that is, in 164 a good solvent. However, in a bad solvent, this effect must be 165 taken into account, considering that the structure and solubility 166 of chitosan is strongly influenced by the pH of the solution.⁴¹ 167

To promote deprotonation of chitosan, in the titration 168 measurements, the pH of the liposome suspension was 169 increased from 4.48 ± 0.01 to 7.10 ± 0.01 . The pH of the 170 original chitosan solution was kept constant at 4.48 ± 0.01 . In 171 this way, when fully (100%) protonated chitosan at pH 4.48 is 172 titrated in liposome suspension with higher pH, some amino 173 groups must deprotonate.

On its own, the deprotonation of chitosan is strongly 175 endothermic, that is, producing strong endothermic titration 176 peaks as shown in Figure 1B. The polymer was titrated in a 177 f1



Figure 1. Molecular structure of deionized/deprotonated chitosan (A). Isothermal titration calorimetry traces (25 °C) for the titration of chitosan solution at concentration of 1 mg/mL (6.13 mM of monomers) at pH 4.48 \pm 0.01 (80 mM acetate buffer) in the same buffer at pH 7.10 \pm 0.01 (B), and for the titration of chitosan-free buffer with pH 4.48 into the corresponding buffer at pH 7.10 (C). The exothermic heat contribution due to buffer mixing is small (less than 10% in magnitude) compared to the endothermic heat of chitosan dilution.

buffer with pH 7.10 in the absence of vesicles. Even very slight 178 differences in pH produce high endothermic signal when the 179 pH is higher than that of the chitosan solution. The reference 180 titration reflecting the contribution of buffer ionization is also 181 shown, see Figure 1C, where the chitosan-free buffer at pH 4.48 182 was injected into the buffer at pH 7.10. The heat signal of this 183 buffer mixing is exothermic and small in magnitude (less than 184 10%) compared to the heat absorbed upon chitosan dilution. 185

t1

186 Titrations in buffers with pH 4.90, 5.30, and 6.00 have shown 187 qualitatively similar results (data not shown).

To understand the nature of the strong endothermic signal 188 189 associated with the dilution of protonated chitosan in buffers 190 with higher pH, we first consider pure dilution of the polymer. 191 In a previous study,³⁵ we have shown that the injection of 192 chitosan solution into the same (native) buffer solution (with 193 identical pH), produces composite peaks. Each injection consists of a small exothermic peak immediately followed by 194 195 an endothermic one, resulting in an overall endothermic signal 196 of only around 2 μ J per injection; see also Figure S1 in the 197 Supporting Information. However, as shown in Figure 1B, the simple dilution of the fully protonated polysaccharide in a 198 solution with increased pH results in strong endothermic effect. 199 The signal is more than 2 orders of magnitude higher; compare 200 with data in Figure S1 in the Supporting Information. This 201 202 behavior must be associated with deprotonation of the amino group of the chitosan monomers to reach charge equilibrium in 203 204 the new solvent. The deprotonation of the polymer is also 205 evidenced from the decrease in the ζ -potential measured in 206 solutions of increasing pH; see Table 1. Further contributions

Table 1. Chitosan Characteristics in the Native Buffer at pH 4.48 \pm 0.01 and after Titration in 80 mM Acetate Buffer of Different pH^a

initial pH	final pH	particle diameter (nm)	ζ-potential (±5 mV)	conductivity (mS/cm)	
	4.48 (native)	87 ± 26	53	3.92	
4.90	4.81	112 ± 36	47	3.86	
5.30	5.10	134 ± 33	42	4.13	
6.00	5.42	477 ± 182	40	4.41	
7.10	5.64	aggregates	40	4.78	

^aThe first column indicates the buffer pH before mixing, and the second column shows the conditions after the titration and at which the DLS and ζ -potential measurements were performed. The errors in the particle diameter indicate standard deviations. The standard deviation in the ζ -potential measurements performed on the same sample was less than 1 mV. In the column title, we have indicated the instrument accuracy as specified by the manufacturer. The monomeric concentration of the original chitosan solution was 6.13 mM (95% DDA, 199 kDa). See text for details.

207 to the heat release in the titration curves may be related to 208 structural changes in the polymer. Indeed, deprotonation of

chitosan can be expected to result in conformational changes, 209 whereby the extended chains may bend and fold adopting a 210 new organization in order to reduce the contact with the polar 211 water molecules. Such structural changes can produce 212 endothermic signal resulting from delocalization of water 213 molecules and deprotonation, both taking place as the 214 conformation of the polymer is changing. Another effect to 215 be considered above a certain degree of deprotonation is the 216 aggregation of the polymer in solutions with higher pH. 217 Evidence for aggregation is provided from dynamic light 218 scattering (DLS); see Table 1. The measured size increases 219 systematically after titration in solutions with increasing pH. 220 Titration in buffer with pH 6.00 leads to a more pronounced 221 size increase, while in solutions of pH 7.10 micrometric 222 particles were detected. 223

Whenever pH values of solutions are adjusted, one typically 224 adds small amounts of strong acid, which may significantly alter 225 the ionic strength of the solution and consequently affect the 226 electrostatic interactions. In the measurements above, we took 227 extra care to carefully adjust the pH to avoid strong variations 228 in the ionic strength. For monitoring purposes, we measured 229 the solutions conductivities to make sure that the ionic strength 230 of the solutions did not vary significantly; see last column in 231 Table 1. Solutions with excessively high deviations in the 232 conductivity, more than around 0.7 mS/cm compared to the 233 average value, were discarded as they would correspond to a 234 change in the ionic strength by more than around 10 mM. 235

When two buffers are mixed in titration, both bear a buffering 236 effect, which leads to a different pH of equilibrium as indicated 237 in the second column of Table 1. Hence, the associated heat 238 absorption is higher in the first injections (Figure 1B), where 239 the pH difference between the mixed solutions is maximal. 240 Subsequently, the signal decreases after each new injection, 241 meaning that the new polymer chains introduced in the buffer 242 with the following injections suffer less deprotonation than the 243 chains from the previous injections, as a consequence of pH 244 reduction from the buffering effect.

To summarize, the heat release in the ITC measurements can 246 be ascribed to the combined effect of different contributions, 247 namely, deprotonation of amino groups, changes in polymer 248 conformation, and polysaccharide aggregation. All these effects 249 are more pronounced when the change in pH is larger. 250

3.2. Interaction of Deprotonated Chitosan with 251 Liposomes. Having characterized the effects associated with 252

Table 2. Results for the ζ -Potential and the Hydrodynamic Diameter of DOPC/DOPG Liposomes (90/10, 3.82 mM total phospholipid concentration in 80 mM acetate buffer) in Buffers of Varied pH at 25 °C after Titration with the 80 mM Acetate Buffer with Constant pH of 4.48 and after Titration with Chitosan Solution (6.13 mM of monomers in 80 mM acetate buffer, pH 4.48)^{*a*}

					titration of buffered chitosan in vehicles			
			titration of buffer in vesicles				degree of degree $(1 - 6)$	eprotonation α) (%)
vesicle pH	pH of added solution	final pH	ζ -potential (±5 mV)	particle diameter (nm)	ζ -potential (±5 mV)	particle diameter (nm)	first injection	final injection
4.90	4.48	4.82	-26	93 ± 3	41	223 ± 14	7	6
5.30	4.48	5.11	-29	103 ± 4	39	239 ± 23	17	11
6.00	4.48	5.43	-21	98 ± 4	38	aggregates	50	21
7.10	4.48	5.64	-24	109 ± 3	39	aggregates	93	30

^{*a*}The errors in the particle diameter indicate standard deviations. The standard deviation in the ζ -potential measurements on the same sample was less than 1 mV. In the column title, we have indicated the instrument accuracy as specified by the manufacturer. The calculated percentage of deprotonation degree (1/ α) of chitosan for each vesicle pH is shown in the last two columns; see text for details.

253 mixing chitosan with buffers at different pH, we now proceed 254 with discussing the interaction of the polymer with the 255 membrane. As a charged model membrane, we have chosen 256 DOPC vesicles containing 10 mol % DOPG (DOPC/DOPG 257 90/10) since these fractions of charged lipid have yielded 258 reasonable ITC signal.³⁵ Furthermore, the pK_a of DOPG is 259 around $3^{42,43}$ which implies that the surface charge of the bare 260 vesicles remains negative after titration of buffer with higher 261 pH. This is confirmed by the ζ -potential measurements given in 263 altered when the liposomes get in contact with the native buffer 264 of the chitosan solutions.

f2

f3

²⁶⁵ Figure 2 shows an example titration of chitosan in a vesicle ²⁶⁶ suspension at pH 5.30. The first few injections produce a



Figure 2. ITC trace (25 °C) for DOPC/DOPG liposomes (90/10, 3.82 mM total phospholipids) at pH 5.30 \pm 0.01 (80 mM acetate buffer) titrated with chitosan solution (1 mg/mL, 6.13 mM of monomers) at pH 4.48 \pm 0.01 (same acetate buffer).

267 composite signal: a positive (upward) endothermic peak 268 immediately followed by a negative (downward) exothermic 269 peak. The magnitude and trend of the exothermic part of the 270 signal is similar to that observed when titrating the same 271 chitosan solution in the vesicle suspension (DOPC/DOPG 90/ 272 10) but at pH 4.48;³⁵ see Figure S2 in the Supporting 273 Information. This suggests that the exothermic part of the 274 signal is associated with the neutralization of negative charges of DOPG by the positive ionized amino groups of chitosan. 275 276 However, now the exothermic effect is quickly suppressed by the endothermic signal associated with chitosan deprotonation. 277 After the third injection, the peaks turn highly positive and the 278 279 endothermic effect predominates until the end of the titration with intensity similar to that of the reference titration (titration 280 of chitosan solution in the corresponding vesicle-free buffer, 281 compare with Figure 1B). This trend was observed for all 282 283 measurements at different pH.

After subtracting the reference measurement from the main 284 titration, that is, deducting the effect associated with chitosan 285 dilution and deprotonation, one obtains the net signal 286 associated with polymer-vesicle interaction. In Figure 3, we 2.87 present the data in terms of integrated heat per injection for 288 every different pH shift. The first injection point is excluded 289 from the data since it is strongly influenced by dilution effects 290 during the pre-equilibration stage of the measurement. The 291 results in Figure 3A are presented in terms of interaction with 292 the accessible lipid considering that chitosan can interact only 293 with the external leaflet of the vesicle membrane because at the 294 295 explored conditions, no bilayer poration has been observed; see 296 accompanying study.³⁶ The data show that the heat released 297 from DOPG neutralization is highly exothermic and the signal 298 increases for higher differences in the pH between the chitosan 299 solution and the vesicle suspension. Above molar ratios of 0.1



Figure 3. (A) Integrated heat per injection versus the molar ratio of chitosan to phospholipids present in the external leaflet of the liposome membrane for the titrations of chitosan solution (1 mg/mL, 6.13 mM monomeric chitosan) at pH 4.48 \pm 0.01 (80 mM acetate buffer) in DOPC/DOPG small liposomes (90/10, 3.82 mM total phospholipids) in 80 mM acetate buffers with different pH (\pm 0.01) as indicated. The heat of chitosan dilution has been subtracted from the data. The inset plot shows the decrease in molar enthalpy of polymer–lipid interaction as a function of pH in the solution (the first data point for pH 4.48 is taken from ref 35); see text for details. The slope of the linear fit is (-7.08 ± 0.71) kJ/mol per pH unit. (B) Molar enthalpy of interaction as a function of the degree of deionization (deprotonation) of the polymer calculated for the first injection of the ITC measurement.

chitosan monomers-to-accessible lipid, the released heat levels 300 out close to zero for all samples, suggesting, once again, 301 neutralization of DOPG on the membrane of the vesicles (note 302 that the molar fraction of DOPG in the membrane is 10%). 303 However, when varying the pH of the liposome suspension, 304 one has to consider the deprotonation of chitosan as an 305 effective process that influences the interaction of the polymer 306 with the vesicle membrane. 307

We did not pursue model fitting of the titration data sets to 308 extract binding constants because of the relatively weak signal 309 and the lack of strong functional dependence in the explored 310 interval of molar ratios. However, we performed the following 311 rough analysis to evaluate the molar enthalpy of binding. The 312 data was extrapolated to zero molar ratio of chitosan monomers 313 to accessible lipid; see dashed curves in Figure 3A. In this 314 regime of excess lipid, one can assume that all injected chitosan 315 fully engages in binding to the membrane. The released heat 316 will be then directly proportional to the molar enthalpy of 317 interaction, ΔH , and given by the intercept. The inset in Figure 318 3 shows the linear variation of the molar enthalpy ΔH_{319} estimated in this way as a function of the pH of the vesicle 320 solutions. The higher the pH, the larger in magnitude the 321 enthalpy of interaction is demonstrating the stronger 322 exothermic effect as a function of increasing pH. 323

Considering the different acidity conditions analyzed in this 324 work, buffers with pH values of 4.90 and 5.30 are still 325 reasonably good solvents for chitosan and when the polymer is 326

327 introduced in these buffers only a small amount of protons 328 must be transferred to the solution. However, taking into 329 account the enthalpy deduced for the first injections in 330 solutions of pH values 4.90 and 5.30 (see inset in Figure 331 3A), it is reasonable to admit that even weak deprotonation of 332 chitosan influences the energy of electrostatic interaction, 333 where the enthalpy changes almost by a factor of 2, from -3.87334 kJ/mol of chitosan monomer in pH 4.90 to -6.60 kJ/mol of 335 chitosan monomer in pH 5.30.

On the other hand, the solutions with higher pH values of 337 6.00 and 7.10 are no longer good solvents for chitosan and 338 deprotonation of the amino groups is largely increased when 339 the polymer is introduced. Chitosan becomes unstable in these 340 buffers and exhibits a critical tendency to form aggregates 341 (Table 1).

Following an approach introduced by Rinaudo et al.,⁴¹ we stimated the degree of deprotonation of chitosan for each pH evaluated in this work using the expression

$$pK_a = pH + \log_{10}[\alpha/(1-\alpha)]$$

345 where pK_a for chitosan is assumed to be 6.0^{41} and α is the 346 degree of protonation, that is, the opposite of the degree of 347 deprotonation $(1 - \alpha)$. For the native chitosan solution with 348 pH 4.48, the degree of protonation is 97%, or equivalently in 349 terms of deprotonation degree $1 - \alpha = 3\%$. These values imply 350 full solubility of the polymer chains. As expected, with 351 increasing pH, the deprotonation degree increases as evidenced 352 in Table 2. At pH 6.00, it reaches 50%, which is considered the 353 limit for chitosan solubilization.⁴¹ The buffer at pH 7.10 is no 354 longer a good solvent for the polysaccharide, since the 355 deprotonation is higher than 90%.

Let us mention the following caveat here. Such deprotonation degrees may be effective only for the first couple of injections of chitosan solution into the buffers with higher pH. The reason for this is because both solutions have a buffering effect, as discussed above, and their mixing produces a solution with final pH at the end of the titration, also shown in Table 2. Thus, following the first injections, deprotonation of chitosan still occurs, but evidently to a lesser extent since the pH in the stitration cell is slowly decreasing with each new injection.

Simultaneously with the polymer deprotonation, the 365 366 neutralization of negative charges on the membrane of the vesicles containing 10% DOPG also occurs during the first 367 368 injections of chitosan solution. We consider an approximated 369 deprotonation degree at this point, as given in the penultimate 370 column of Table 2. When injected into the vesicle suspension, 371 the polymer binds onto the negatively charged membranes (note that the membrane surface charge is independent of pH; 372 373 see Table 2) and an exothermic signal from electrostatic 374 interaction is released indicating stabilization of the system. The 375 higher the difference in pH, and thus the instability of chitosan 376 due to deprotonation, the higher the enthalpic contribution 377 upon binding will be as the polymer acquires stability on the vesicle surface, as shown in Figure 3B. Thus, it is evidenced that 378 the state of lower energy for the system is when chitosan is 379 adsorbed on the membrane of the vesicles instead of being free 380 381 in solution and in a bad solvent.

After the neutralization of DOPG, basically only endothermic heat of deprotonation of the polymer chains characterizes the set system. Once again, the decrease in the endothermic signal for set ach subsequent injection of chitosan solution observed in Figures 1B and 2 is related to the systematic reduction of pH in the ITC reaction cell due to the addition of increasing amounts of chitosan buffer at pH 4.48. As shown in Table 2, the final pH 388 in the cell is around 5 depending on the initial pH of the 389 vesicles buffer. In this way, the decrease in the endothermic 390 signal also shows the decrease of chitosan deprotonation as a 391 function of pH. 392

Let us now consider the results for the ζ -potential and 393 particle size in the deprotonation experiments. The data given 394 in Table 2 evidence that simple titration of chitosan-free buffer 395 with pH 4.48 into vesicle suspensions with increasing pH values 396 produces neither a change in the size nor in the ζ -potential of 397 the bare vesicles. All vesicles remain with around 100 nm of 398 hydrodynamic diameter and surface charge close to -25 mV. 399 After titration with chitosan solution, the vesicles in solutions of 400 pH 4.90 and 5.30 behave similarly exhibiting a substantial 401 increase in size and ζ -potential, suggesting chitosan adsorption. 402 However, the vesicles in solutions at pH 6.00 and 7.10 show the 403 presence of microaggregates after titration with chitosan, but no 404 further increase in the ζ -potential. Thus, the adsorption of 405 chitosan on the membrane of small liposomes may be 406 influenced also by the degree of deprotonation of the polymer. 407

To summarize, the adsorption of polymer chains on the 408 liposomes is irreversible and alters the vesicle membrane 409 characteristics. The mechanical properties of the membranes 410 are also significantly altered as explored in the subsequent study 411 employing giant vesicles.³⁶ Indeed, the aggregation observed 412 here by means of the DLS measurements may be responsible 413 for increasing the membrane apparent roughness as visualized 414 with microscopy observations on giant vesicles incubated in 415 chitosan solutions. The change in surface charge from negative 416 for the bare vesicles to positive for the chitosomes might be 417 responsible for the observed adhesion and rupture of giant 418 vesicle in contact with glass surfaces.³⁶

3.3. Polymer Reorganization. Previously, we have 420 proposed a model to explain the organization of the 421 electrostatically driven binding of chitosan onto liposomes 422 with varied surface charge.³⁵ Here, we discuss a model to 423 address the reorganization of the polymer under the effect of 424 deprotonation. As illustrated in Figure 4A, for pH values lower 425 f4



Figure 4. Schematic representation of chitosome structures. (A) Positively charged chains of chitosan are attracted by the slightly negative surface charge of the small liposome in buffer with $pH \leq 5.30$. (B) Aggregates of chitosomes formed after the partial deprotonation of chitosan in buffer with $pH \geq 6.00$. The polymer chains bind to the membrane of the slightly negatively charged liposomes and mediate adhesion based on hydrophobic attraction between the deionized/ deprotonated parts of chitosan. The overall surface charge becomes positive; see text for details.

and around 5.30 where the deprotonation of chitosan is low, 426 the polymer chains are attracted by the slightly negatively 427 charged DOPC/DOPG 90/10 vesicles and adsorption must 428 occur in a similar fashion as previously described for these 429 structures. As a consequence of the adsorption of the bulky 430 polymer chains, the chitosome size increases and the ζ - 431 potential becomes highly positive, preventing aggregation 432 (Table 2).

In solutions with pH 6.00 and above, the deprotonation of 434 435 chitosan is increased (Table 2) and the polymer is no longer 436 stable in the buffer. The negative charges on the vesicles are 437 available in the same buffer and chitosan is strongly attracted, 438 leading to stronger adsorption as shown by the increased 439 exothermic heat release when comparing the ITC data collected 440 for different pH values; see Figure 3B. The stronger attraction 441 which has to occur fast (and mainly in the first injections) may 442 lead to lower organization of the polymer over the membrane, 443 leading to chitosome aggregates once again, as discussed 444 before.³⁵ However, since the surface charge of the DOPC/ 445 DOPG 90/10 vesicles is only slightly negative, the final ζ -446 potential turns now positive (Figure 4B), contrary to what was 447 observed for vesicles with higher DOPG fractions.³⁵ In 448 addition, as a consequence of deprotonation of chitosan, 449 hydrophobic domains are produced on the chains presumably 450 leading to changes in the conformation of the polymer as a 451 result of intramolecular and intermolecular hydrophobic 452 interactions.²⁷ Furthermore, the deionized/deprotonated seg-453 ments of the chains must have looser binding over the 454 membrane of the vesicles, since electrostatic interaction with 455 these segments is no longer possible. Thus, for $pH \ge 6.00$, one 456 could expect that upon adsorption to the liposome surface the 457 polymer chains attain conformations, which differ from those at 458 lower pH as illustrated in Figure 4. Differently from the 459 previous report,³⁵ where the aggregation may be produced by 460 looser chains attracted by the strong negative charge of 461 neighboring vesicles, in the present study, the net charge is no 462 longer negative as shown by the increase in ζ -potential (Table 463 2), suggesting that now the aggregation is caused mainly by 464 hydrophobic interactions between the deprotonated loops of 465 the looser chains.

To summarize, the deprotonation of chitosan at high pH 467 leads to polymer reorganization that strongly influences the 468 interaction and adsorption of the chains on the liposome 469 membranes. For the intermediate pH values, between 5.30 and 470 6.00, the chitosan behavior in solution is difficult to evaluate 471 and more effort has to be made in this direction. In this pH 472 range, the polymer solubility changes significantly as the degree 473 of deprotonation increases and intramolecular interactions 474 appear to influence the conformation of the polymer. The 475 polymer behavior may also be influenced by the distribution of 476 remaining acetyl groups as well as by the distribution of 477 molecular weight, as chitosan behavior in such solutions is far 478 from ideal.⁴⁰

In order to better understand the polysaccharide—membrane interaction and enlighten the physical characteristics associated with the encounter of chitosan with phospholipid membranes, we addressed the effect of the polymer on the membrane rigidity and pore formation in giant unilamellar vesicles with similar lipid composition.³⁶

4. CONCLUDING REMARKS

485 The interaction between chitosan and phospholipid vesicles 486 may be enhanced by means of altering compositional and 487 environmental conditions. Here, we demonstrated the use of 488 chitosan deprotonation as a tool to modulate the polymer 489 adsorption to the membrane. The deprotonation of chitosan 490 was promoted with titration of the protonated polymer into 491 solutions with increasing pHs and the process was found highly 492 endothermic. This endothermic effect increases with pH 493 showing that the chitosan structure and behavior are strongly 494 dependent on the media acidity. Partially deionized/deprotonated chains still exhibit affinity to the negative surface charges 495 of the membrane. Further deprotonation of the polymer seems 496 not to lead to weaker affinity. Indeed, when bound, chitosan 497 reaches higher stability compared to when it remains free in a 498 poorly protonating solvent, a fact that was shown by the 499 exothermic signal in the ITC titration. However, it has to be 500 stressed that, once again, stronger adsorption results in lower 501 organization of the polymer chains over the membrane. This 502 lower organization, in addition to conformational changes of 503 the chains due to deprotonation, also promotes aggregation of 504 the chitosome structures. Since the final surface charge on 505 chitosomes is highly positive in this case, hydrophobic 506 interactions between deprotonated looser loops of chitosan 507 segments must be related to the aggregation process.

The findings of the present study highlight the importance of 509 degree of chitosan protonation in research areas where chitosan 510 is employed as a macromolecule for biological and biomedical 511 applications. More specifically, the mucoadhesive properties of 512 liposomes coated with chitosan, intended as specific drug 513 delivery systems, may be improved with the knowledge about 514 the degree of chitosan protonation. The polymer deprotonation 515 will influence the degree of coverage in these coated liposomes 516 and can be used to modulate it. In a following study,³⁶ we 517 demonstrate that, on giant vesicles, this coverage is very 518 heterogeneous and depends on the preparation protocol 519 employed.

ASSOCIATED CONTENT					
Supporting Information	522				
ITC data. This material is available free of charge via the					
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Notes					
The authors declare no competing financial interest.					
	535				

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