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Abstract: Immobilization of  $\beta$ -galactosidase ( $\beta$ -gal) into porous materials might afford to supported biocatalysts for the hydrolysis of diary products or to food additives for lactose intolerant people. Activity and stability of the loaded material generally depend on the interactions between the enzyme and the support, as well as on the pore size. Herein, Raman spectroscopy was used to evaluate the specific adsorption the enzyme into meso-macroporous silica materials, containing interconnected mesopores of 9 nm and macropores of 200 nm. Non-porous silica was used as reference material to determine the Raman fingerprint of physisorbed enzyme in the absence of any confinement. While the  $\beta$ -gal physisorbed on the surface of non-porous silica material exhibits the same Raman spectra as the free enzyme, the enzyme physisorbed onto meso-macroporous materials show frequency displacements of characteristic amide groups as a function of initial concentration of the feed enzymatic solution. In fact, at low initial concentration in enzyme, no shifts of the amides were recorded on Raman spectra as compared with free enzyme, indicating a preferential physisorption into macropores. By increasing the enzyme concentration, the frequency of Amide I was shifted to lower values, suggesting thus a confinement into mesopores. Finally, the enzyme concentration effect can be demonstrated by the increment of the amide band intensity in the range of 1700-1500  $\text{cm}^{-1}$  as the amount of adsorbed enzyme increases. Thus, the textural properties of silica materials seem to be the key factor in the enzyme adsorption.

## HIGHLIGHTS

1.  $\beta$ -gal physisorbed on the surface of non-porous silica material or into macropores exhibits the same Raman fingerprint as the free enzyme
2. The characteristic amide bands of  $\beta$ -gal physisorbed into mesopores are red shifted compared to free enzyme
3. Spectral Raman fingerprint probes the confinement of  $\beta$ -galactosidase into meso-macroporous silica.

## Probing the confinement of $\beta$ -galactosidase into meso-macroporous silica by Raman spectroscopy

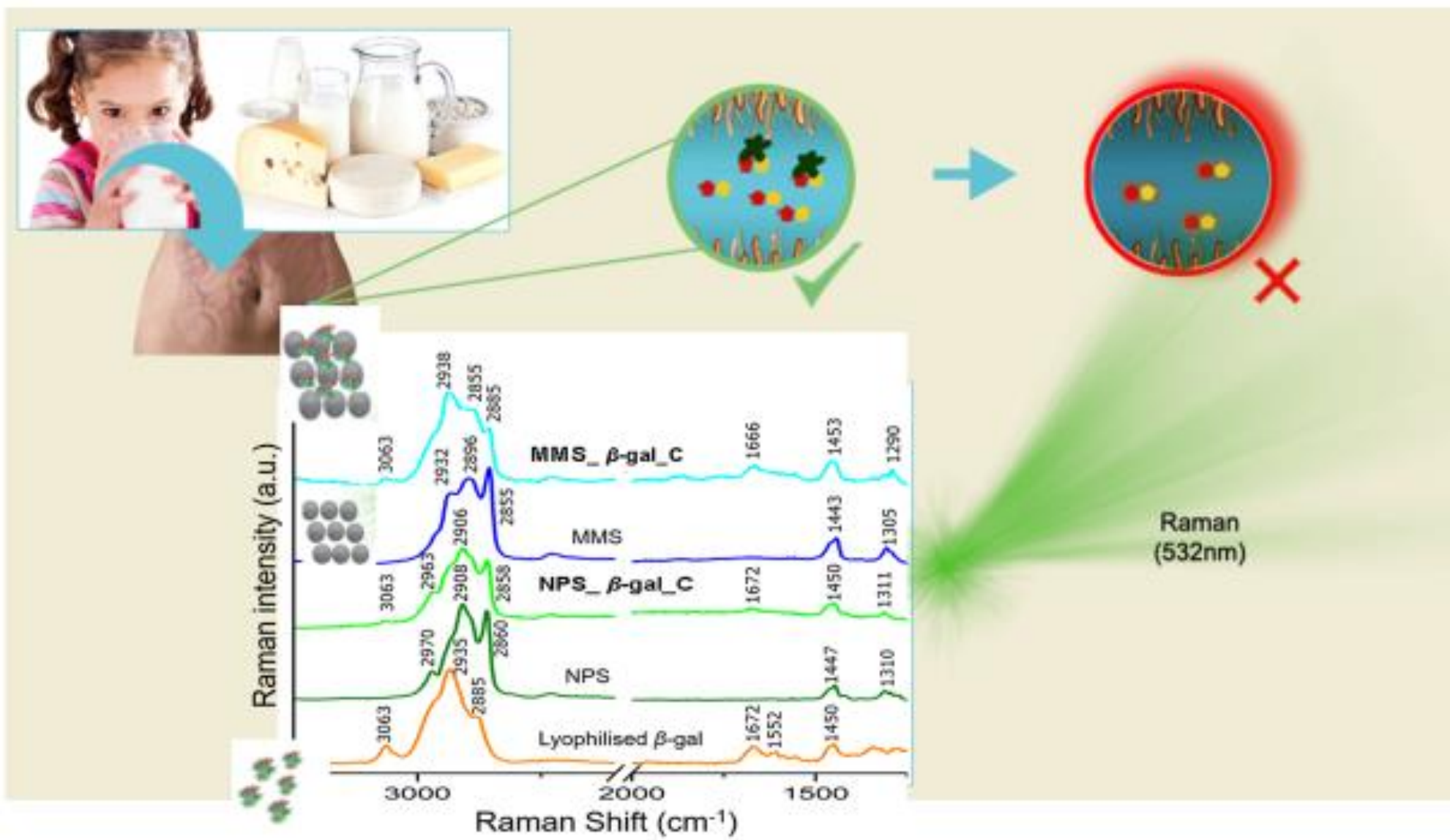
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# 1 Probing the confinement of $\beta$ -galactosidase into meso-macro 2 porous silica by Raman spectroscopy

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44 adsorbed enzyme increases. Thus, the textural properties of silica materials seem to be the key  
45 factor in the enzyme adsorption.

46

47 **Keywords:**  $\beta$ -galactosidase, biocatalysts, meso-macroporous silica, enzyme-support  
48 interactions, Raman spectroscopy.

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## 51        **1. Introduction**

52        Immobilization of biomolecules (i.e. proteins, DNA) into nanostructured materials is extensive  
53        area of research. This is particularly relevant for designing sustainable biocatalysts and  
54        biosensors, allowing for multiple reuses and for the protection of biomolecules from  
55        denaturing external factors or operational parameters such as temperature, pH or reactive  
56        species [1-5]. Many organic or inorganic porous supports such as polymers, carbon nanotubes  
57        or particles, metal-oxides, glass or porous silica materials have been investigated for  
58        biomolecules immobilization, especially due to their capacity of loading as well as mechanical  
59        and chemical stability [4]. Among those supports, ordered porous silica materials are the most  
60        widely chosen for the following main reasons: the tunability of the surface chemical functions  
61        and the control of the pore size to fit with the biomolecule diameter and to increase the specific  
62        surface of the support [6-9].

63        Enzyme-supported silica materials already found industrial and biotechnological applications,  
64        including in food industry. In this field, one challenge still lies in finding solutions for lactose  
65        intolerant consumers, such as for the production of low-lactose products. This could be achieved  
66        by using lactase enzymes, *e.g.*  $\beta$ -galactosidase ( $\beta$ -gal). The immobilization of  $\beta$ -gal into porous  
67        materials might afford to supported biocatalysts for the hydrolysis of dairy products or to food  
68        additives for lactose intolerant people. The enzyme immobilization on a support material is a  
69        good solution to protect the enzyme from external conditions as pH and temperature [10-11].  
70        Among several materials (such chitosan, agarose, and cellulose) [12-13], the porous silica  
71        materials have been widely used as support materials for the immobilization of  $\beta$ -galactosidase  
72        [12-20], mainly due to their large interacting surface and the precise tuning of the pore sizes.  
73        Several strategies have been applied for controlled delivery of  $\beta$ -gal, using both physical and  
74        covalent immobilization of the enzyme. In brief, the attachment of a lactose derivative as  
75        gatekeeper on the surface of mesoporous silica nanoparticles has resulted in successful enzyme-  
76        responsive intracellular controlled delivery of biologically active enzyme  $\beta$ -gal into cells, which  
77        provides promising delivery systems for large therapeutic proteins [6, 7, 17]. Bernal et al.  
78        reported that the three-dimensional structure of the  $\beta$ -gal was not significantly affected during

79 the physical immobilization process on an hierarchical porous silica with a stable large porosity  
80 (10-40 nm and 0.07-20  $\mu\text{m}$  diameter ranges) [14]. Instead of that,  $\beta$ -gal presented a selective  
81 physical adsorption on hierarchical meso-macroporous silica materials (average pore size 9 and  
82 200 nm, respectively), arising from the oligomeric complexation of the enzyme (as  
83 monomer/dimer/tetramer), which affected to the catalytic activity [18]. That enzyme had an  
84 improvement of mechanical and thermal stability by multipoint covalent immobilization in  
85 hierarchical macro-mesoporous silica (average mesoporous diameter 20 nm) [15].

86 The activity and stability of the loaded material generally depend on the interactions between  
87 the enzyme and the support, as well as on the pore size. To gather further insights on those  
88 specific interactions, vibrational spectroscopy (e.g. infrared and Raman spectroscopy) appear as  
89 the most powerful as they provide information about molecular vibrations. Whereas IR spectra  
90 depends on polar moment and tend to emphasize asymmetric vibrations of polar groups (e.g.  
91 OH, C-H, N-H, C=O, etc.), Raman spectroscopy depends on polarizability changes and tends to  
92 emphasize symmetric vibrations of non-polar groups such as C=C and C-C stretches, and  
93 aromatic ring breathing vibrations [21]. Therefore, the frequencies of those molecular vibrations  
94 depend not only on the atom masses and on their geometric arrangement, but also on the  
95 strength of their chemical bonds and interactions. In addition, these techniques present several  
96 attractive features such as being generally fast, non-destructive, and non-invasive.

97 Usually, the presence of enzymes on different supports has been demonstrated by the  
98 appearance of amide bands in the IR spectrum [22-26]. However, only few of those studies are  
99 addressing the fine interactions between the enzyme and the support that lead to different band  
100 frequencies on the IR spectrum [22, 25]. Some authors have justified the interactions by the  
101 frequency shift of the characteristics bands of the functional groups of the enzymes (mainly  
102 amide) simply by an increase of the band intensity. In addition, different studies have also been  
103 carried out by Raman spectroscopy for different enzymes/supports yielding information about  
104 chemical modification occurring by functionalization, or the study of the enzyme structural  
105 changes induced by the isotopically replacement of  $\text{H}_2\text{O}$  solvent by  $\text{D}_2\text{O}$  [27-31].



106 Indeed, very few studies on the physico-chemical interactions investigated by IR have dealt  
107 with the adsorption of  $\beta$ -gal on different support materials [18, 22, 24], whereas Raman  
108 spectroscopy has not yet been used specifically to identify and characterize  $\beta$ -gal  
109 immobilization. The goal of this work is to show the capabilities of Raman spectroscopy to  
110 characterize the specific adsorption of  $\beta$ -gal onto meso-macroporous silica substrates through  
111 the analysis of interactions between the enzyme and the silica support. It concerns more  
112 particularly the evaluation of the specific adsorption of the enzyme into meso-macroporous  
113 silica materials, containing interconnected mesopores of 9 nm and macropores of 200 nm. A  
114 non-porous silica material was used as reference to determine the Raman fingerprint of  
115 physisorbed enzyme in the absence of any confinement.

116

## 117 **2. Experimental Sections**

### 118 **2.1. Materials**

119 Two different silica supports were used: commercially available non-porous nanoparticles from  
120 Nanocomposites company (NPS, 97 nm diameter), and a porous, homemade material described  
121 by Pasc et al [32], which is a dual meso-macroporous silica (MMS). In brief, the synthesis  
122 protocol involved the preparation of a colloidal suspension of solid lipid nanoparticles of n-  
123 hexadecylpalmitate dispersed in a micellar solution of Pluronic P123 surfactant. A hybrid  
124 organo-silicate material was obtained by the hydrolysis and the polycondensation of  
125 tetramethoxysilane (TMOS) around to the colloidal template. The meso and macro porosities  
126 were released by Soxhlet extraction.

127 The  $\beta$ -gal enzyme solution, extracted from *Kluyveromyces lactis* source, was a gift from Chr.  
128 Hansen Company (Copenhagen, Denmark). The stock solution provided is 80% pure enzyme  
129 (determined by SDS-PAGE analysis) and has a concentration of 40 mg/mL of protein in PEM  
130 buffer solution (phosphate, ethylenediaminetetraacetic acid (EDTA), and magnesium, pH 6.6)  
131 and glycerol (1:1) weight ratio. The enzyme activity of stock solution was measured at 5200  
132 NLU/g. It was determined spectrophotometrically from its ability to convert *o*-nitrophenyl- $\beta$ -D-

133 galactopyranoside (ONPG) into galactose and *o*-nitrophenol (ONP). Assay of  $\beta$ -gal activity was  
134 described in reference [11]. One NLU is the quantity of enzyme that releases 1.3  $\mu$ mol of *o*-  
135 nitrophenol per min at 30°C and pH 6.5 [33]. All the buffer reagents were purchased from  
136 Sigma-Aldrich (Spain).

## 137 **2.2. Immobilization of $\beta$ -gal onto silica supports**

138 The enzyme immobilization was performed on two types of silica supports (porous and non-  
139 porous) through physical adsorption, by mixing the enzyme solution and the silica material in  
140 the PEM buffer (pH 6.6), under agitation for 48 hours at room temperature. The amount of silica  
141 was fixed (25 mg) and the  $\beta$ -gal concentration was varied according to the concentration of the  
142 feed solution denoted as C/x, where x indicates the dilution factor. Thus, the C/x concentrations  
143 ranged from C (40 mg/ml) to C/2, C/4, C/6, C/10 and C/20. After immobilization, samples were  
144 washed 3 times with buffer solution, by centrifugation at 4000 rpm. Then, samples were dried at  
145 room temperature, and a white powder was obtained. Finally, the samples were kept at room  
146 temperature.

147 The protein loading in the respective silica materials were determined using a UVIKON 943  
148 UV-Vis Kontron spectrophotometer, thought their absorption wavelength at 280 nm. To do so,  
149 the absorbance of each supernatant recovered after the washing step was measured. Table 1  
150 summarizes the amount of  $\beta$ -gal adsorbed on the MMS and was calculated as the difference  
151 between the initial amount of protein in the respective adsorption solution and the residual  
152 amount of the  $\beta$ -gal lost in the supernatant. The samples are denoted from now on as, for  
153 example, “MMS\_  $\beta$ -gal\_C/2” meaning dilution of factor 2 of the stock enzyme solution  
154 adsorbed on the meso-macroporous silica support.

155 **Table 1.** Amount of  $\beta$ -gal adsorbed on MMS per surface area of MMS, for the different concentrations  
156 of initial  $\beta$ -gal solutions used for immobilization.

Sample	Initial $\beta$ -gal concentration (mg/mL)	Adsorbed $\beta$ -gal (mg/ m <sup>2</sup> )
MMS_ $\beta$ -gal_C	40	16
MMS_ $\beta$ -gal_C/2	20	10
MMS_ $\beta$ -gal_C/4	10	8.0
MMS_ $\beta$ -gal_C/6	6.7	6.9

MMS_β-gal_C/10	4.0	5.2
MMS_β-gal_C C/20	2.0	5.2

157 Concerning the sample NPS\_β-gal\_C, the adsorbed enzyme was 0.117 mg/m<sup>2</sup> of silica NPS.  
 158 That corresponds to two orders of magnitude lower than the analogous sample for the MMS  
 159 support.

160 The immobilized enzyme was active for both silica systems and for all the different feed  
 161 concentrations. As reported in previous work [18], the specific activity of the enzyme  
 162 physisorbed in the macropores was two times higher (about 30-35 U/g) than that of the enzyme  
 163 entrapped into mesopores (below 15 U/g), where 1 U corresponding to the amount of enzyme  
 164 catalyzing the conversion of 1 μmol of ONPG per minute at 25 °C. For the NPS\_β-gal system,  
 165 the activity was inspected qualitatively, verifying the yellow color after convert ONPG into  
 166 galactose and ONP, when reaction was carried out by the same assay methodology than for the  
 167 hierarchical porous silica system.

## 168 **2.3. Characterization**

### 169 **2.3.1. Morphological properties**

170 The morphology and the porosity at the bare silica materials were observed by Transmission  
 171 (TEM) and Scanning (SEM) Electron Microscopy. The TEM pictures were carried out with a  
 172 Philips CM200 at 200 kV and the Image processor software used was GATAN. The MMS  
 173 powder was first ground and then suspended in ethanol (Sigma-Aldrich, France) by sonication.  
 174 A drop of the dispersion was spread out on the TEM carbon lacey grid and dried at room  
 175 temperature before observation. The SEM micrographs were obtained with a Zeiss DSM 950  
 176 scanning electron microscope using an acceleration voltage of 15kV. The NPS powder was  
 177 lyophilized before to air-disperse on carbon grid followed by metal coating with a Polaron E-  
 178 5400 equipment. The images were acquired from randomly chosen location and analysed with  
 179 the scanning probe image processor software (Quartz PCI).

### 180 **2.3.2. Structural properties**

181 The elementary organic analysis of silica materials was performed using a LECO CHNS-932  
182 Elementary Chemical Analyzer.

183 Fourier Transform Infrared spectroscopy (FTIR) measurements were performed using a Thermo  
184 Scientific FT-IR Nicolet IS 10 (Waltham, MA, USA), equipped with an attenuated total  
185 reflectance (ATR) sampling accessory with a diamond crystal. The version 6.0 of Omnic  
186 software from Thermo Scientific was used for instrument control and spectra acquisition. Each  
187 absorbance spectrum was recorded from 4000 to 400  $\text{cm}^{-1}$  with a resolution of 4  $\text{cm}^{-1}$  and the  
188 signal averaged over 32 scans. The spectra of four independent spots of each sample were  
189 collected, and then, a mean spectrum was calculated and finally normalized between 0 and 1.

190 Raman spectra were collected using a Thermo Scientific confocal DXR Raman Microscope  
191 controlled by the Thermo Scientific OMNIC for dispersive Raman 8.3.103 software. A laser  
192 emitting at 532 nm with 8 mW and a confocal slit size of 50  $\mu\text{m}$  were used. The microscope was  
193 set to 50x magnification under bright field illumination. Raman spectra of all samples were  
194 collected in the range 3450-500  $\text{cm}^{-1}$ . The spectral resolution of the used instrument  $\pm 2 \text{ cm}^{-1}$ .  
195 The number of scans was optimized for the different samples in order to obtain the best signal-  
196 to-noise ratio (always within the range 5-50 scans), while the optimum exposure time for each  
197 scan was fixed to 10 seconds. Fluorescence correction (polynomial order 6 of OMNIC  
198 fluorescence correction) and smoothing (5 points, polynomial order 2) were applied. All Raman  
199 spectra were finally normalized between 0 and 1 in order to facilitate the comparison between  
200 the recorded spectra according the enzyme content and the silica support type.

201 No sample treatment was performed to bare silica materials, but the  $\beta$ -gal standard solution was  
202 lyophilised before Raman analysis. Both silica supports were microscopically different.  
203 Contrary to the homogeneous spherically shaped non-porous silica, the home synthesized meso-  
204 macroporous silica showed a big heterogeneous surface. For both supports, focusing was an  
205 extremely difficult task because the samples were composed of a white powder, assembled in  
206 different layers inducing different surface depths. To avoid this problem, Raman spectra were  
207 collected on a very small powder grain, but the laser spot size (about 1.2  $\mu\text{m}$ ) always resulted  
208 smaller than the particle size. In addition, microscope glass slides were previously covered with

209 aluminium foil to avoid the spectral interferences caused by glass when measuring the particles  
210 of silica material.

211 Due to the heterogeneous adsorption (*i.e.* heterogeneous distribution of  $\beta$ -gal on silica supports),  
212 there were some spots, in particular for low enzyme contents, where the amide bands were  
213 intense and dominated the Raman spectra, and some other spots, where it was only possible to  
214 detect the bands of the silica material. To overcome this limitation, spectra from five random  
215 selected spots of the same sample were acquired at room temperature to record and averaged  
216 signal intensity.

217 The graphs presented in this paper have been plotted using the OriginPro® 9.1 software  
218 (Northampton, MA, USA).

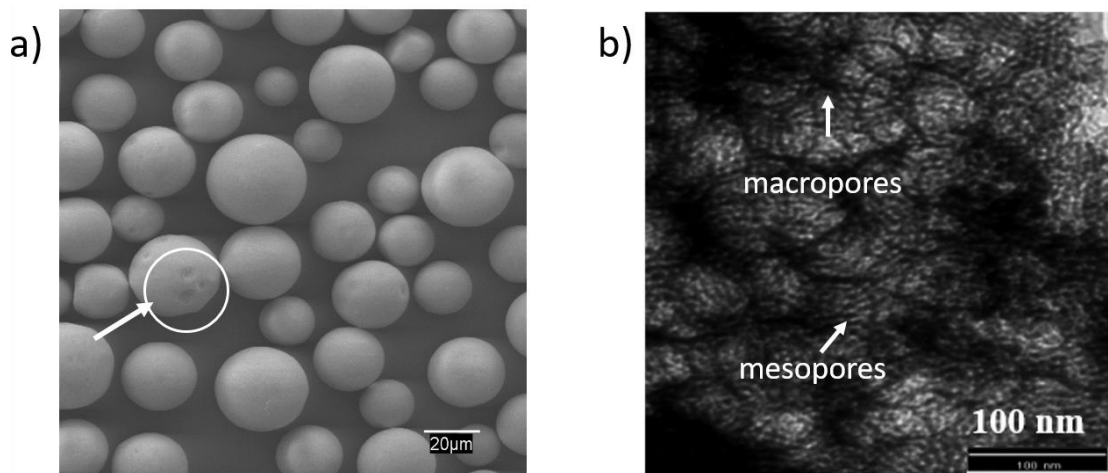
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### 220 **3. Results and discussion**

221 The  $\beta$ -gal was adsorbed in both types of silica materials as described in the experimental  
222 section. We proposed two different particles morphologies in order to inspect the capability of  
223 Raman spectroscopy to evaluate the strength of the expected different interaction between the  
224 enzyme and the two silica supports. The main difference between the systems is the  
225 confinement of the enzyme in the meso and macropores of the MMS materials.

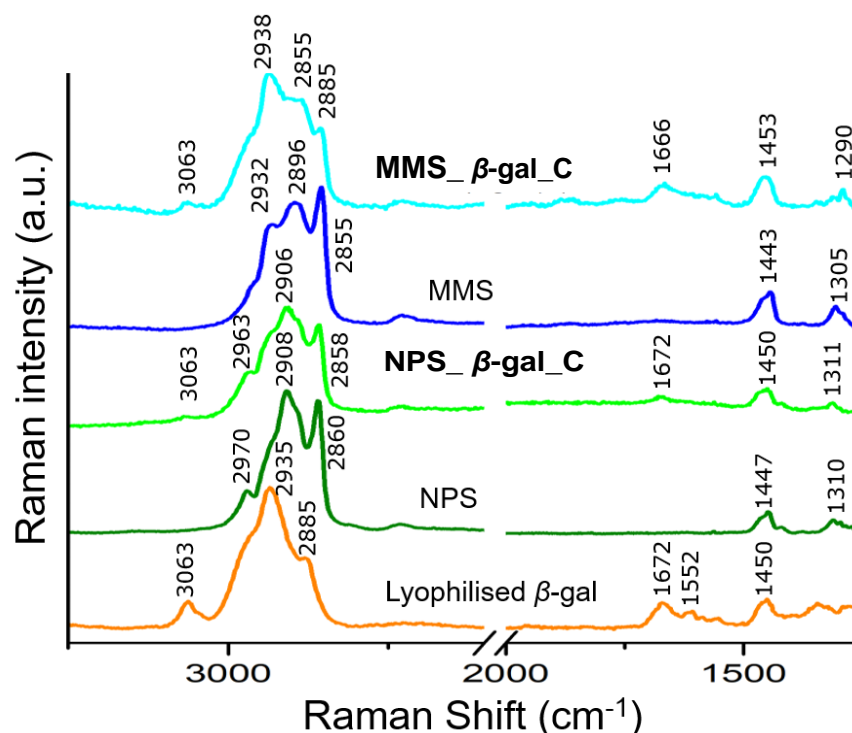
226 Firstly, the morphology of the meso-macroporous silica and the non-porous silica was analysed  
227 by TEM and SEM, respectively, as shown in Figure 1. The non-porous silica are spherical  
228 particles of about 30  $\mu\text{m}$  that display sporadically micro-craters of about 2  $\mu\text{m}$  on the surface  
229 according to SEM photomicrographs (Fig. 1A). The MMS material support shows a dual meso-  
230 macroporosity of 9 nm and 100 nm, respectively (Fig. 1B). As determined from nitrogen  
231 sorption measurements [18], bare MMS material had a specific surface area of 660  $\text{m}^2 \text{g}^{-1}$  and a  
232 pore volume of 1.23  $\text{cm}^3 \text{g}^{-1}$ . The material exhibited a type IV isotherm, characteristic of a  
233 mesoporous material, with an average pore size of 9 nm, and a steep increase of the adsorbed  
234 volume at  $p/p_0 > 0.9$ , confirming the presence of macropores. The  $\beta$ -gal is a homo-oligomer of  
235 four subunits with dimensions of 15.1 nm  $\times$  17.1 nm  $\times$  10.7 nm, as inferred from the crystal

236 structure, leading to an averaged hydrodynamic diameter of about 14 nm. On the other hand, the  
237 dimer and the monomer can be contained in boxes of dimensions  $11.9 \text{ nm} \times 15.6 \text{ nm} \times 7.2 \text{ nm}$ ,  
238 and  $7.2 \text{ nm} \times 11.7 \text{ nm} \times 6.3 \text{ nm}$ , respectively. On the basis of this data, it was reasonably  
239 hypothesized that monomers and dimers can be adsorbed into the mesopores, while the  
240 tetramers can be physisorbed onto the macropores. As evidenced in previous work, the  
241 preference for being in dimer or tetramer form mainly depended on the enzyme concentration  
242 [18].



243 **Figure 1.** (A) SEM images of NPS particles where the arrows and the circles point the macropores and  
244 (B) TEM images of MMS.  
245  
246

247 The Raman spectra of both bare silica supports were recorded and compared with the signal of  
248 lyophilized free enzyme and loaded MMS and NPS silica materials at the highest concentration  
249 (MMS\_β-gal\_C and NPS\_β-gal\_C) as presented on Figure 2.



250 **Figure 2:** Raman spectra of NPS and MMS before and after  $\beta$ -gal immobilization (loaded at the  
 251 maximum concentration, C, from the enzyme stock solutions). Raman spectra of lyophilized  $\beta$ -gal  
 252 free enzyme is also provided for comparison purposes.  
 253

254 The spectra of both bare silica supports (MMS and NPS) present some bands in the range 3000-  
 255 2500  $\text{cm}^{-1}$ , which are assigned to the stretching vibrations of C-H groups. Particularly, they are  
 256 related to the compounds used in the synthesis process of these silica materials, *i.e.* some traces  
 257 of the surfactants and/or lipids used in the synthesis as templates can remain even after  
 258 calcination [34]. This is in agreement with the thermogravimetric analysis of the MMS bare  
 259 silica support that contains almost 21 wt % organics [18]. However, the Raman shift of the C-H  
 260 bands of the non-porous silica material (2970, 2908, 2860  $\text{cm}^{-1}$ ), and the relative intensity of  
 261 them, are different to the ones of the porous silica (2932, 2896, 2855  $\text{cm}^{-1}$ ). No information was  
 262 available about the reagents used in the manufacture of the commercial non-porous silica  
 263 support, but the elementary analyses support the presence of some organics too. The nitrogen  
 264 and sulphur atoms are not present in the bare materials, which allows us to rule out the presence  
 265 of bacterial contamination or microorganisms according to elemental analysis data.

266 Figure 2 also shows the spectrum of the lyophilised  $\beta$ -gal which exhibits a band at 3063  $\text{cm}^{-1}$   
 267 due to N-H stretching vibrations, characteristic of amides and commonly designated as amide B  
 268 band [28, 29]. There is also an overlap of several intense bands within the range 3000-2850  $\text{cm}^{-1}$

269 (maxima at 2935 cm<sup>-1</sup> and 2885 cm<sup>-1</sup>), characteristic of C-H stretching vibrations, which belong  
 270 to the backbone of the different amino acids that constitute  $\beta$ -gal. The bands located within the  
 271 1700-1500 cm<sup>-1</sup> range were assigned to the overlap of C=O stretching and N-H bending  
 272 vibrations, representative of Amide I and Amide II bands, respectively. This band assignment  
 273 was previously discussed by Baranska et al [35]. Furthermore,  $\beta$ -gal displays bands in the 1470-  
 274 1250 cm<sup>-1</sup> range corresponding to C-H bending vibrations [35-36]. The Table 2 summarizes the  
 275 main Raman bands of the non-porous silica support (NPS), the meso-macroporous silica support  
 276 (MMS) and  $\beta$ -gal enzyme with their assignment to the respective characteristic molecular  
 277 vibrations [35-38].

278

279

280

281 **Table 2.** Main Raman bands from the collected spectra (NPS, MMS and free  $\beta$ -gal enzyme) and their  
 282 assignment with the fundamental vibrations, according to the literature [35-38]. (str: stretching, bend:  
 283 bending, as: asymmetric, s: symmetric).

Sample	Raman Shift (cm <sup>-1</sup> )	Molecular vibration
NPS	2970 2908 2860 1447 1310	str(as) CH <sub>3</sub> str CH <sub>2</sub> str(s) CH <sub>3</sub> bend C-H bend C-H
MMS	2932 2896 2855 1443 1305	str(as) CH <sub>3</sub> str CH <sub>2</sub> str(s) CH <sub>3</sub> bend C-H bend C-H
$\beta$ -gal enzyme	3063 2935 2885 1672 1552 1470-1250	str N-H (Amide B) str C-H str C-H str C=O (Amide I) str C-N and bend N-H (Amide II) bend C-H

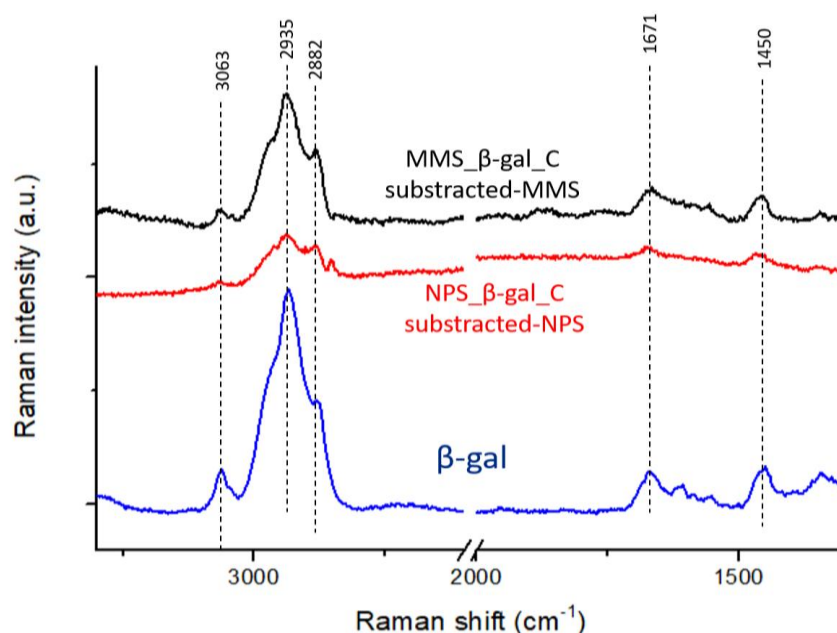
284

285 With regards to the spectrum of the MMS materials after  $\beta$ -gal immobilization (loaded at the  
 286 maximum concentration, C), the appearance of the bands corresponding to Amide I, Amide II  
 287 and Amide B proved the presence of the enzyme on the silica support (Figure 2). It should be



288 noted that Amide I band suffered a slight displacement to  $1666\text{ cm}^{-1}$  when comparing with the  
289 free enzyme ( $1672\text{ cm}^{-1}$ ) (Raman uncertainty  $\pm 2\text{ cm}^{-1}$ ). These amide bands were also observed  
290 in the non-porous silica support after  $\beta$ -gal immobilization, though they did not display any shift  
291 and they were relatively less intense than for the porous silica reacting with the same  $\beta$ -gal feed  
292 solution. The intensity of the amide bands is reasonably smaller since the amount of adsorbed  
293 enzyme is much lower (two orders of magnitude) for the NPS support, at the same enzyme feed  
294 solution, C (Table 1).

295 For better understanding and clearer visualization of the Raman bands of  $\beta$ -gal in the  
296 MMS/NPS supports after  $\beta$ -gal immobilization, the background provided by each support was  
297 subtracted (Figure 3). This way, the characteristic Raman bands of  $\beta$ -gal are clearly  
298 recognizable in both supports (not only amide I band at  $1671\text{ cm}^{-1}$ , but also the C-H stretching  
299 bands located within  $3000\text{--}2850\text{ cm}^{-1}$ , which were previously overlapped in Figure 2 by C-H  
300 bands of the supports). Furthermore, after subtracting the background support, it can be also  
301 noticed in Figure 3 that the amount of adsorbed enzyme on MMS support is larger than the  
302 amount adsorbed on NPS support (even though the same enzyme concentration was added to  
303 both supports). As described above, this result evidences the higher adsorption of  $\beta$ -gal on  
304 MMS in comparison to NPS.



305 **Figure 3:** Raman subtracted spectra of NPS and MMS after  $\beta$ -gal immobilization. Upper spectrum is the  
306 result of subtracting MMS spectrum to MMS-  $\beta$ -gal spectrum (of Figure 2). Second spectrum is the result  
307

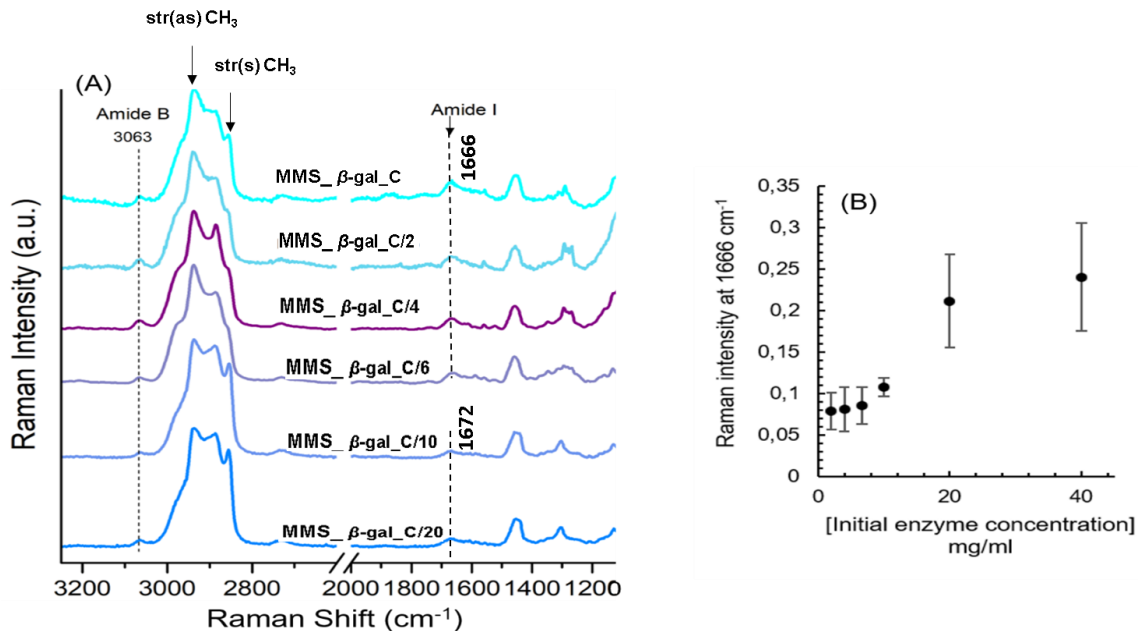
308 of subtracting NPS spectrum to NPS-  $\beta$ -gal spectrum (of Figure 2). Raman spectra of lyophilized  $\beta$ -gal  
309 free enzyme is also provided for comparison purposes.

310 In order to examine the strength of the  $\beta$ -gal-silica interactions, additional washings were  
311 performed for both silica supports after enzyme immobilization as previously explained in the  
312 experimental section. In the case of the  $\beta$ -gal adsorbed on the porous silica support, the  
313 supernatant of the first additional washing did not show the presence of protein. On the  
314 contrary, for the non-porous silica samples, the supernatant of each additional washing (up to a  
315 total of fifteen) still contained released protein, as determined by UV absorption as previously  
316 described. This fact supports the outcome that  $\beta$ -gal poorly interacts with the non-porous silica  
317 support in contrast to the porous silica support.

318 Therefore, it appears clearly that the pores are favouring the interactions between the enzyme  
319 and the meso-macroporous silica by physical interactions (Van der Waals, hydrophobic and/or  
320 electrostatic interaction) leading thus to the entrapment of  $\beta$ -gal. The  $\beta$ -gal presents an  
321 isoelectric point of 5.1 while silica has a pI of 2-3. The immobilization process was performed  
322 at pH 6.6 to ensure the enzyme stability. At that pH, both the enzyme and the silica support are  
323 negatively charged. Previously, it was reported that there is a selective adsorption in the meso-  
324 macroporous, arising from the oligomeric complexation of the enzyme  
325 (monomer/dimer/tetramer) [18]. As previously explained, monomers or dimers (whose size < 9  
326 nm) can go into the mesopores (ca. 9 nm size), while the tetramers (ca. 14 nm) can be only  
327 physisorbed onto the macropores.

328 The dimers interfaces involve a large number of accessible positively charged residues [18] that,  
329 in the presence of negatively charged silanol groups, favour electrostatic enzyme-silica  
330 interactions into the mesopores. In addition, the profile of Raman spectrum of MMS\_ $\beta$ -gal\_C,  
331 with respect to the bare material (MMS) spectrum, changes notably in the 3000-2840  $\text{cm}^{-1}$   
332 range; there is a change in the intensity ratio of the characteristic str(as)  $\text{CH}_3$ /str(s)  $\text{CH}_3$  bands  
333 (see Fig. 2 and Table 2). As this fact does not occur for the adsorption of the enzyme on the  
334 NPS, which also contains organic residues, then, the hydrophobic interactions between the

335 organic residues of the silica materials and the significant proportion of hydrophobic-type  
 336 interfaces of the dimers [18] must be due to the confinement into the pores.  
 337 Then, the sensibility of Raman spectroscopy was inspected for this meso-macroporous support  
 338 by studying different concentrations of adsorbed  $\beta$ -gal. These spectra are shown in Figure 4 A.

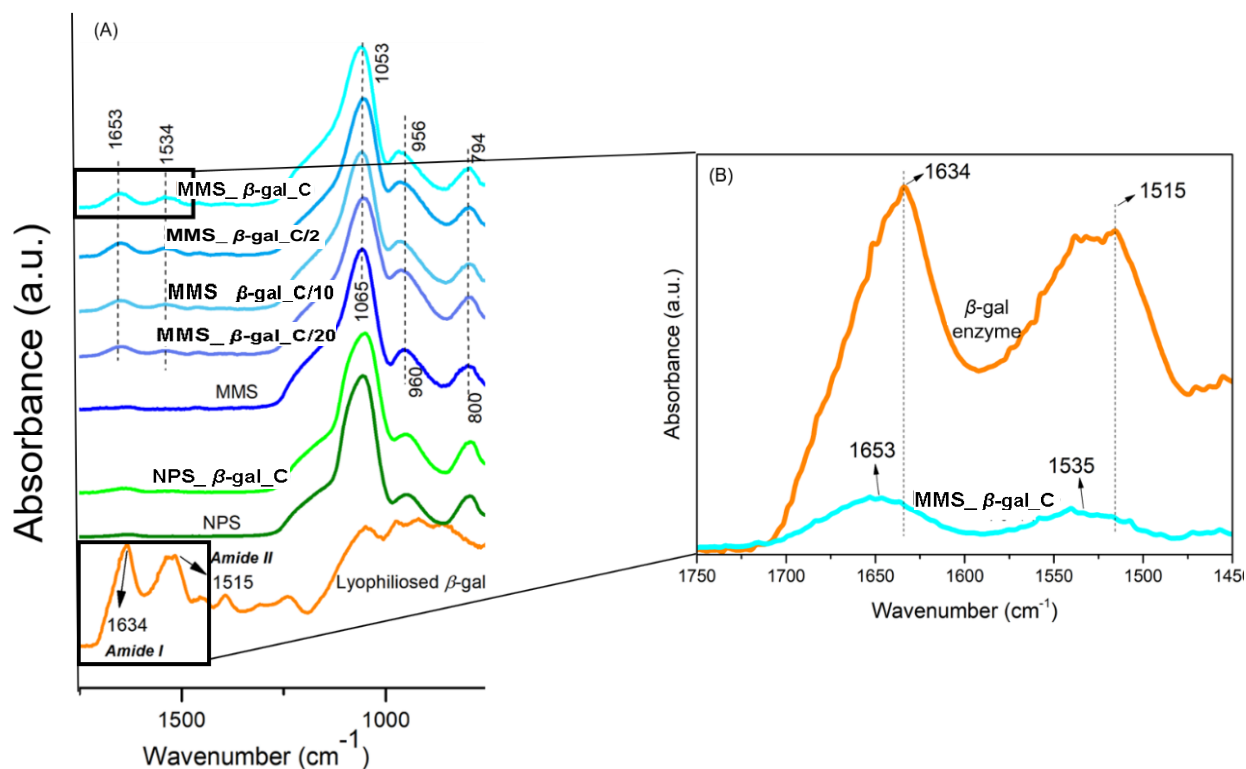


339 **Figure 4.** (A) Raman spectra of meso-macroporous (MMS) silica support after  $\beta$ -gal immobilization, for  
 340 different feed solutions at C, C/2, C/4, C/6, C/10 and C/20 (corresponding adsorbed enzyme is given in  
 341 Table 1); (B) Raman intensity at  $1666\text{ cm}^{-1}$  as a function of the enzyme concentration in the feed solution  
 342 (denoted as initial enzyme concentration). The same trend is followed as a function of the adsorbed  
 343 enzyme.  
 344

345 The increase of the adsorbed enzyme amount may be followed by the intensity's increment of  
 346 the bands located at  $1666\text{ cm}^{-1}$  and  $3063\text{ cm}^{-1}$ , assigned to Amide I and Amide B, respectively.  
 347 In addition, the variation of enzyme concentrations led to a change in the relative intensity of  
 348 the bands between  $2770\text{--}3000\text{ cm}^{-1}$  due to the respective C-H vibrations of the enzyme and the  
 349 support. For the samples containing high enzyme concentrations (C, C/2 and C/4), the band due  
 350 to the contribution of C-H stretching vibrations from the backbone of the amino acids of  $\beta$ -gal  
 351 ( $2935\text{ cm}^{-1}$ ) was the most intense. However, for the samples containing low enzyme  
 352 concentrations (C/10 and C/20), the C-H stretching vibrations from the porous silica support  
 353 ( $2896$  and  $2855\text{ cm}^{-1}$ ) were dominant over the band from  $\beta$ -gal. The spectra also exhibited a  
 354 slight decrease in the relative intensity of the Amide B band for lower enzyme concentrations.

355 Although the change in the spectra profiles could be attributed to the progressively increase of  
356 enzyme content, there is not a linear effect with the loading degree of the enzyme. Particularly,  
357 as displayed in Figure 4 B, the Raman intensity of Amide I band increased with the enzyme  
358 concentration of the feed solution, following a sigmoidal function. The inflection point seems to  
359 be located around 15 mg/mL of enzyme feed solution. The same trend is followed as a function  
360 of the adsorbed  $\beta$ -gal (in mg/m<sup>2</sup> MMS). This result is in concordance to our previous study [18],  
361 where the physisorption mechanism of the enzyme into the meso-macroporous silica materials  
362 depended on the concentration of the enzyme feed solution and on the pore size of the support.  
363 In fact, at low initial concentration in enzyme (MMS\_ $\beta$ -gal\_C/20 and MMS\_ $\beta$ -gal\_C/10), no  
364 shifts of the amides were recorded on Raman spectra as compared with free enzyme (at 1672  
365 cm<sup>-1</sup>), indicating a preferential physisorption into macropores. By increasing the enzyme  
366 concentration, the frequency of Amide I was shifted to lower values (at 1666 cm<sup>-1</sup>), suggesting  
367 thus a confinement into mesopores. The selective physisorption into the mesoporous may arise  
368 from the electrostatic interaction of dimer and silanol groups, compared to the tetramer  
369 complexation at higher concentrated feed solutions that guides to a preferred adsorption of the  
370 enzyme onto the macropores. Higher enzymes concentrations lead to the aggregation of  
371 enzymes that block the diffusion of the enzyme dimers in mesopores [18]. The lower uploading  
372 rate, observed at high concentration of enzymes according to TGA or ATR-FTIR analysis [18],  
373 results in the *plateau* attained for the Raman intensity (Fig 3B). It should be mention that laser  
374 reach enzyme into the mesopores, since the silica wall thickness is lower than 6 nm [18]; i.e.,  
375 concentration of the enzyme affect similarly the Raman intensity if the enzyme is located into  
376 pores or onto external MMS surface. Consequently, both of the red shift of the characteristic  
377 amide bands of  $\beta$ -gal physisorbed into mesopores vs free enzyme, and the sigmoidal increase of  
378 intensity as a function of loading degree, highlight the impact of interactions beyond just an  
379 enzyme concentration effect and the confinement of  $\beta$ -gal into the meso-macoporous silica.

380 IR spectroscopy was used in this study as a complementary technique to Raman spectroscopy.  
381 Figure 5 displays different IR spectra for an increment of enzyme concentration during the  
382 immobilization process for the porous and the non-porous support material.



383 **Figure 5.** (A) ATR-FTIR spectra of  $\beta$ -gal free enzyme, porous (MMS) and non-porous (NPS) silica  
 384 support before and after  $\beta$ -gal immobilization, for different enzyme concentrations (C, C/2, C/10 and  
 385 C/20 (C=40 mg/ml) for  $\beta$ -gal on PS and C for  $\beta$ -gal on NPS). (B) Displacement bands of  $\beta$ -gal free  
 386 enzyme in comparison with  $\beta$ -gal immobilization on the porous silica (NPS).  
 387

388 Figure 5 clearly shows that the intensity of the amide bands is increasing with the  
 389 immobilisation rate. As described for the Raman experiments, comparing at the same enzyme  
 390 feed solution (C, C/2, C/10 and C/20), the intensity of the Amide band I is higher for  $\beta$ -gal  
 391 adsorbed on porous silica (MMS) than on non-porous silica (NPS) samples. The fact is also in  
 392 agreement with the lower enzyme adsorption onto the NPS particles compared to the MMS  
 393 material (see Table 1). Regarding samples of  $\beta$ -gal adsorbed on meso-macroporous silica, the  
 394 band assigned to Si-O-Si stretching vibration at  $1065\text{ cm}^{-1}$  in the spectrum of the bare porous  
 395 silica, shifted to  $1053\text{ cm}^{-1}$  after  $\beta$ -gal immobilization. The band at  $960\text{ cm}^{-1}$  assigned to the Si-  
 396 OH bending vibration, was also displaced to  $956\text{ cm}^{-1}$ . Another band at  $800\text{ cm}^{-1}$  assigned to Si-  
 397 O-C stretching vibrations suffered a displacement to  $794\text{ cm}^{-1}$  after enzyme adsorption on the  
 398 porous support. All those vibrations that involved the silica material have lower energy, and the  
 399 chemical bonds are certainly less interacting with the enzyme.

400 Moreover, the centre of the maxima of the Amide I and Amide II bands of the free  $\beta$ -gal  
401 enzyme shift from  $1634\text{ cm}^{-1}$  to  $1653\text{ cm}^{-1}$  and  $1515\text{ cm}^{-1}$  to  $1535\text{ cm}^{-1}$ , respectively, for the  
402 enzyme-silica samples (see inset Fig. 4). The amide I is sensitive to particular secondary  
403 structural conformation and consists of many overlapping component bands that represents  
404 different structures: purely  $\alpha$ -helices,  $\beta$ -sheets, turns and unordered or irregular structures. The  
405 maximum band at  $1634\text{ cm}^{-1}$  corresponds to  $\beta$ -sheet structures, while  $1653\text{ cm}^{-1}$  is associated to  
406 a more random coil contribution [39]. As the wavelength range of the broad band's contour  
407 doesn't change after the adsorption of the enzyme, the shift of the maxima are related to the  
408 change of integrated intensities areas of its individual underlying band subcomponents, i.e.,  
409 conformational changes of the adsorbed enzyme.

410 Remarkably, no spectral shifts or broadening were observed neither for the enzyme bands  
411 adsorbed on the non-porous material nor for the silica material vibrations. This result  
412 demonstrates again the weaker interaction in the loaded  $\beta$ -gal enzyme onto the non-porous  
413 support in comparison to the porous meso-macroporous support as previously evidenced by  
414 Raman. Thus, the porosity seems to be the key factor in the enzyme entrapment and thus  
415 adsorption.

416

417

418

419

#### 420 **4. Conclusions**

421 In this study, the capability of Raman spectroscopy to prove the adsorption of  $\beta$ -gal on meso-  
422 macroporous silica support was demonstrated by the shifts occurring for the amide bands of the  
423  $\beta$ -gal spectra. In addition, a different profile of Raman spectrum of MMS\_ $\beta$ -gal\_C sample with  
424 respect to the bare material (MMS) spectrum, in the  $3000\text{-}2840\text{ cm}^{-1}$  range, was observed. For  
425 that reason, it is expected that the amide groups are mainly involved in the interaction that  
426 occurred between the enzyme and silica, but also in the hydrophobic forces between the organic

427 residues of the silica material and the aminoacid backbone of the enzyme, probably favoured by  
428 the enzyme confinement into the porous.

429 Thus, the morphology of the silica material seems to be the key factor in the enzyme adsorption.  
430 The physically enzyme adsorption is strongly favoured by the enzyme entrapment into the  
431 pores. While the  $\beta$ -gal physisorbed on the surface of non-porous silica material exhibits the  
432 same Raman spectra as the free enzyme, the enzyme physisorbed onto meso-macroporous  
433 materials show frequency displacements of characteristic amide groups as a function of initial  
434 concentration of the feed enzymatic solution. In fact, at low initial concentration in enzyme, no  
435 shifts of the amides were recorded on Raman spectra as compared with free enzyme, indicating  
436 a preferential physisorption into macropores. In addition, almost all  $\beta$ -gal was released in the  
437 supernatant recovered from the non-porous samples during washing steps, contrary to the  
438 porous samples. All these facts together evidence a stronger interaction between  $\beta$ -gal and the  
439 meso-macroporous silica support than with the NPS material.

440 Finally, it was observed that the intensity of the Raman amide bands bears a sigmoidal  
441 increment by increasing the content of the adsorbed enzyme. The enzyme complexation was  
442 determined in our previous theoretical studies [18]. Further, chemometric analysis could help to  
443 inspect the variability of samples with different amount of adsorbed  $\beta$ -gal on different silica  
444 supports by vibrational spectroscopy.

445

446

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- 562

Figure 1

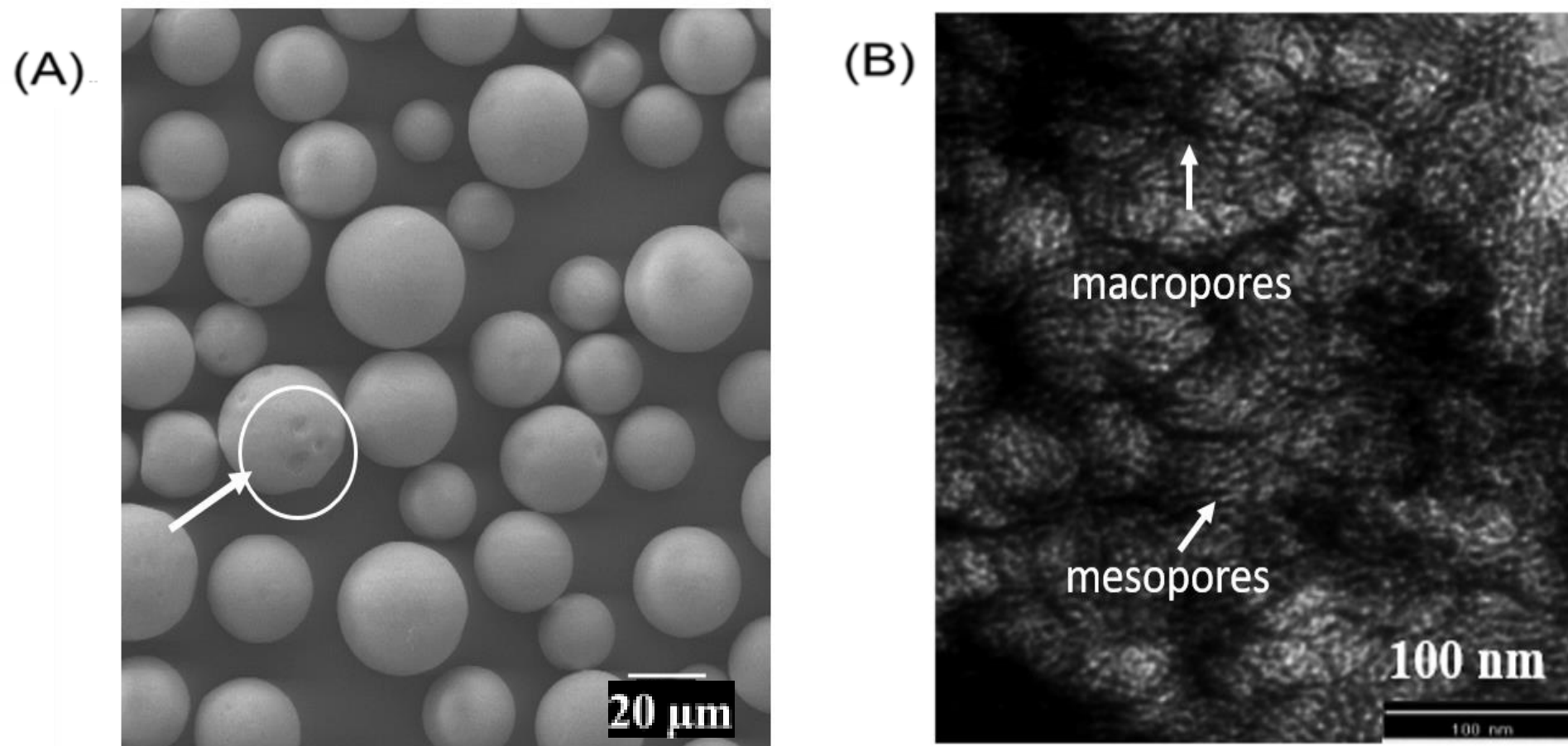


FIGURE 1

Figure 2

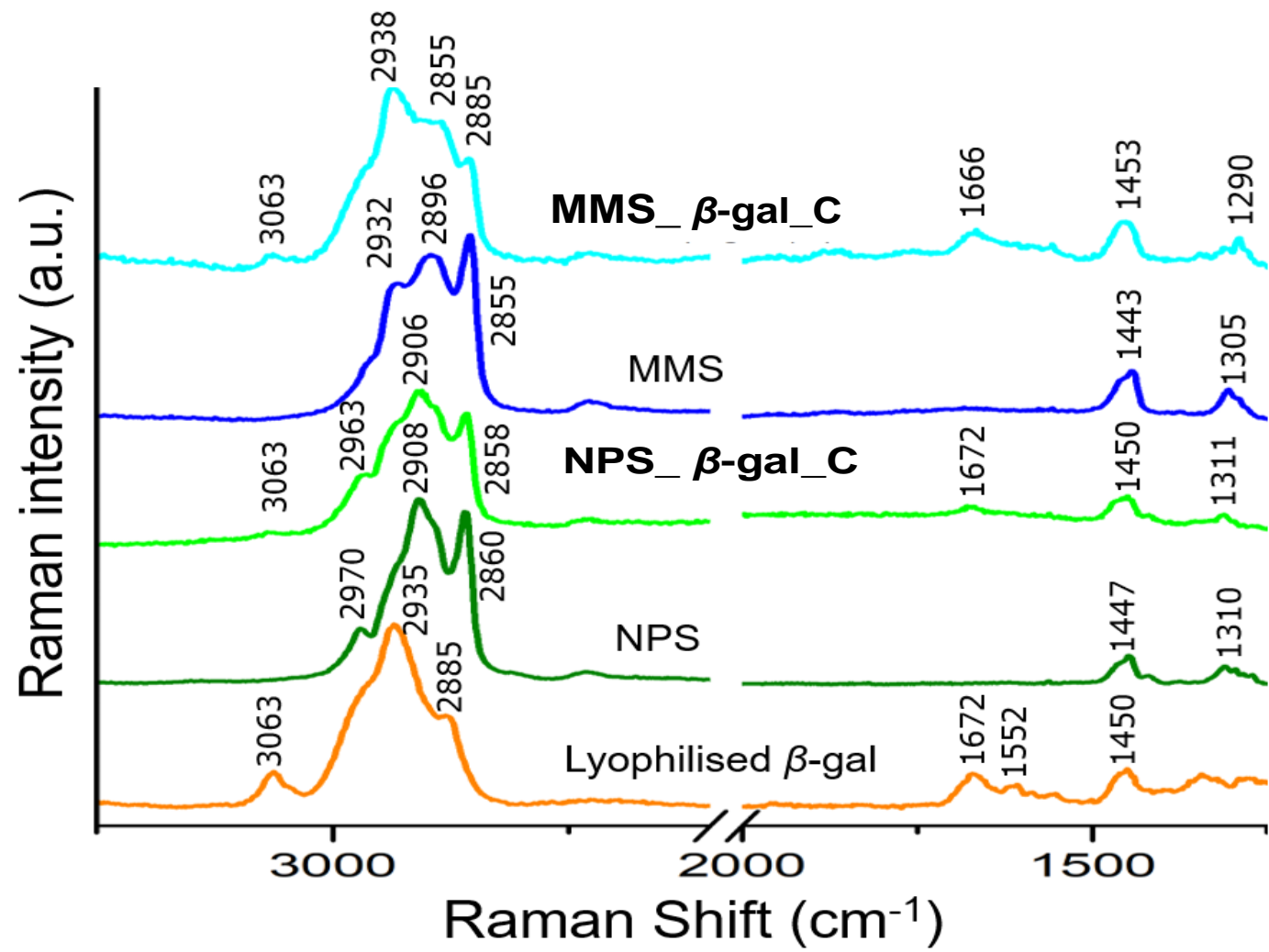


FIGURE 2

Figure 3

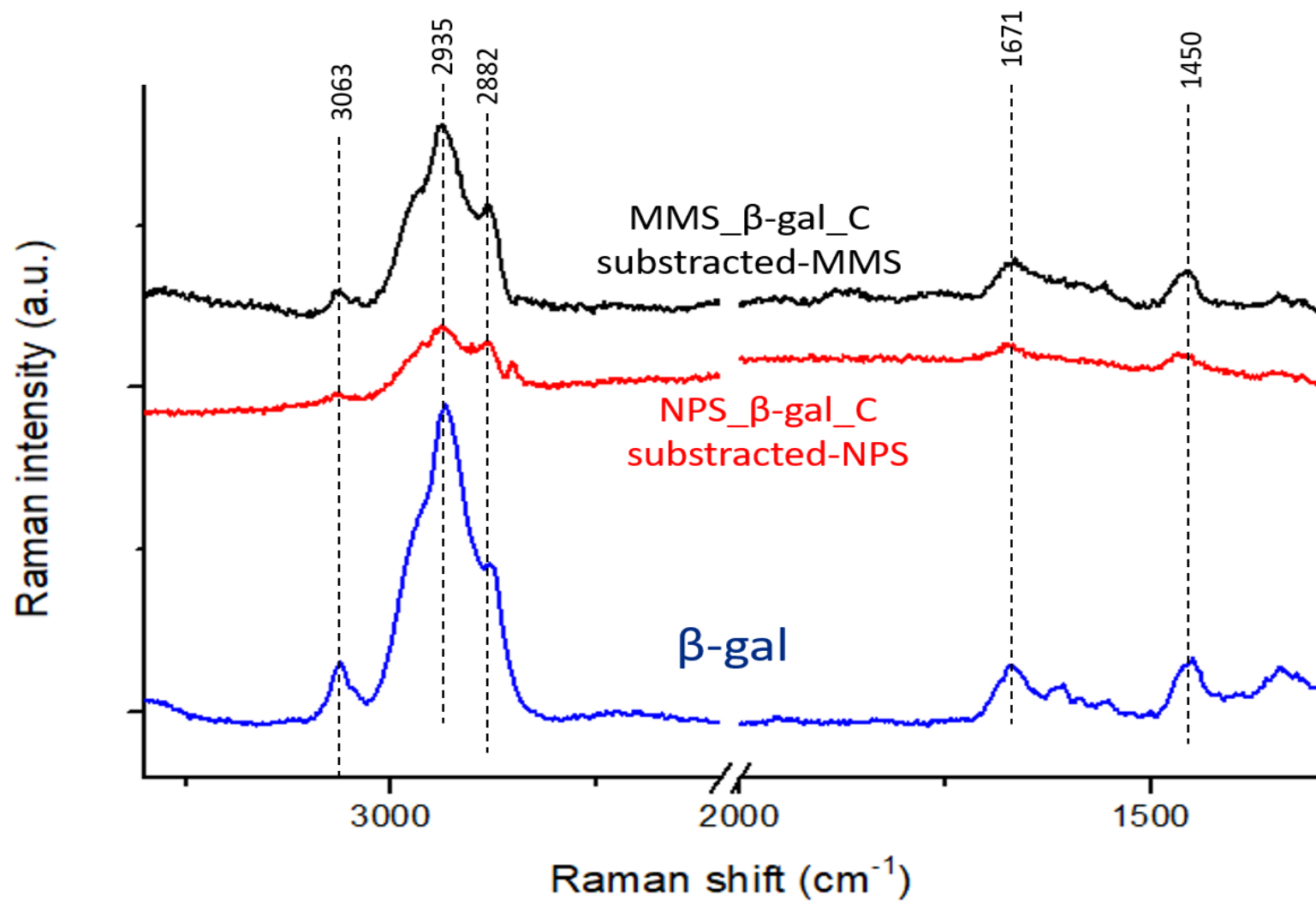


FIGURE 3

Figure 4

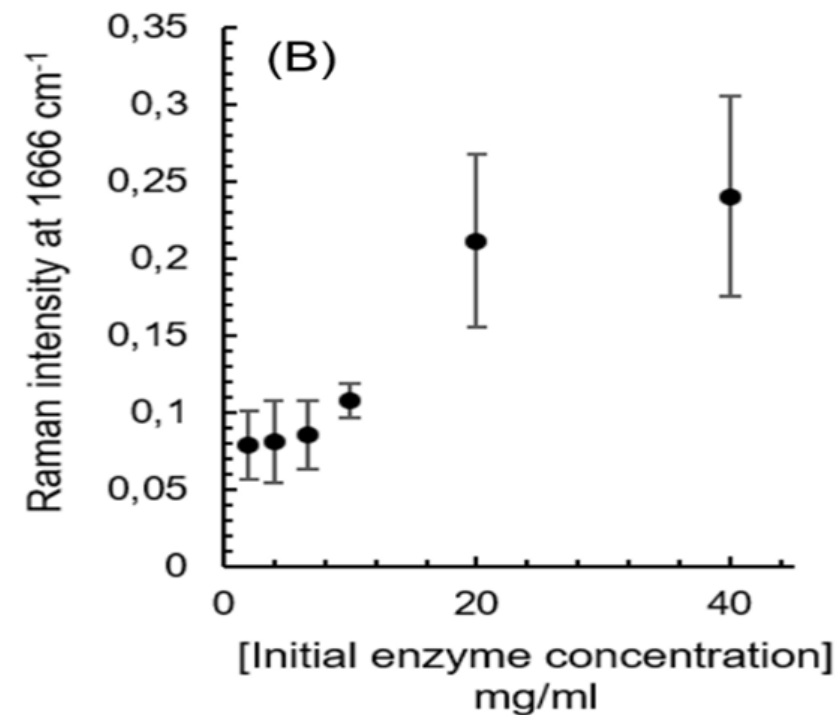
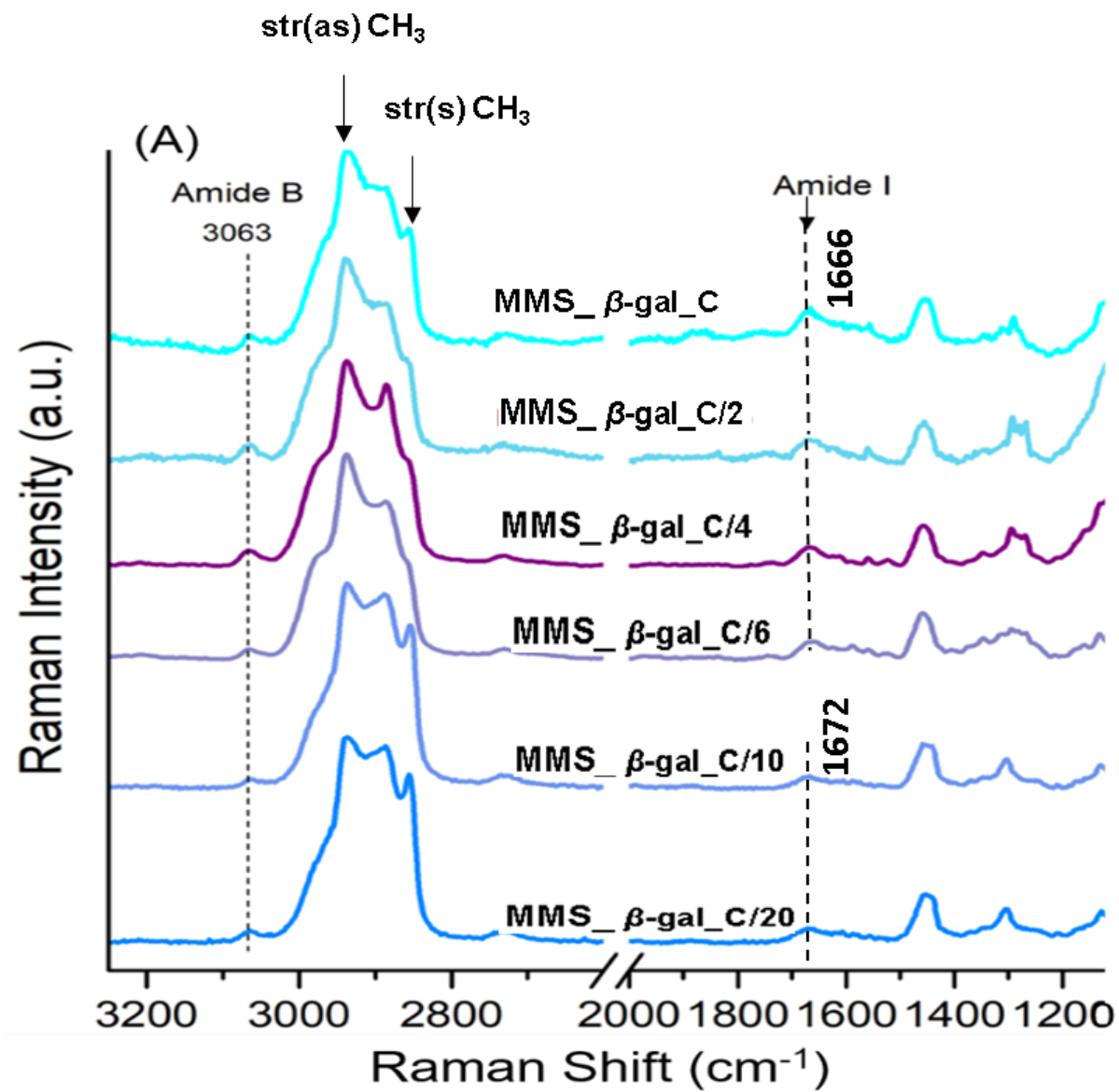


FIGURE 4

Figure 5

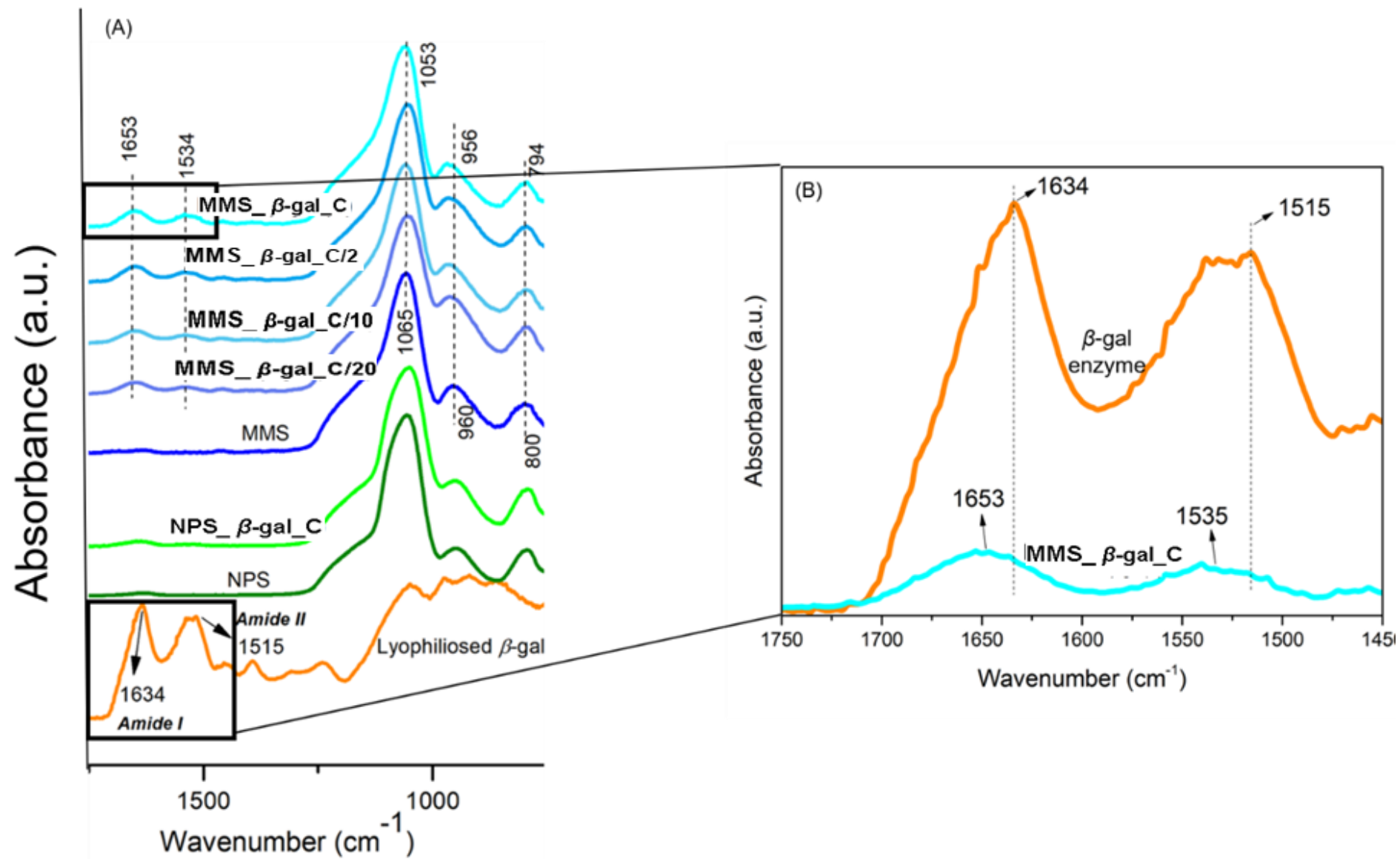


FIGURE 5



**Table 1.** Amount of  $\beta$ -gal adsorbed on MMS per surface area of MMS, for the different concentrations of initial  $\beta$ -gal solutions used for immobilization.

Sample	Initial $\beta$ -gal concentration (mg/mL)	Adsorbed $\beta$ -gal (mg/ m <sup>2</sup> )
MMS_ $\beta$ -gal_C	40	16
MMS_ $\beta$ -gal_C /2	20	10
MMS_ $\beta$ -gal_C /4	10	8.0
MMS_ $\beta$ -gal_C /6	6.7	6.9
MMS_ $\beta$ -gal_C /10	4.0	5.2
MMS_ $\beta$ -gal_C C/20	2.0	5.2

## Probing the confinement of $\beta$ -galactosidase into meso-macroporous silica by Raman spectroscopy

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**Table 2.** Main Raman bands from the collected spectra (NPS, MMS and free  $\beta$ -gal enzyme) and their assignment with the fundamental vibrations, according to the literature [17-21]. (str: stretching, bend: bending, as: asymmetric, s: symmetric).

Sample	Raman Shift ( $\text{cm}^{-1}$ )	Molecular vibration
NPS	2970	str(as) $\text{CH}_3$
	2908	str $\text{CH}_2$
	2860	str(s) $\text{CH}_3$
	1447	bend C-H
	1310	bend C-H
MMS	2932	str(as) $\text{CH}_3$
	2896	str $\text{CH}_2$
	2855	str(s) $\text{CH}_3$
	1443	bend C-H
	1305	bend C-H
$\beta$ -gal enzyme	3063	str N-H (Amide B)
	2935	str C-H
	2885	str C-H
	1672	str C=O (Amide I)
	1552	str C-N and bend N-H (Amide II)
	1470-1250	bend C-H

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