

17 **Abstract**

18

19 The production of new bioactive oligosaccharides is currently garnering much
20 attention for their potential use as functional ingredients. This work addresses the
21 enzymatic synthesis and NMR structural characterization of 2- α -D-glucofuranosyl-
22 lactose derived from sucrose:lactose and sucrose:cheese whey permeates mixtures by
23 using a *Leuconostoc mesenteroides* B-512F dextransucrase. The effect of synthesis
24 conditions, including concentration of substrates, molar ratio of donor/acceptor, enzyme
25 concentration, reaction time and temperature, on the formation of transfer products is
26 evaluated. Results indicated that cheese whey permeate is a suitable material for the
27 synthesis of 2- α -D-glucofuranosyl-lactose, giving rise to yields around 50% (in weight
28 respect to the initial amount of lactose) under the optimum reaction conditions.
29 According to its structure, this trisaccharide is an excellent candidate for a new prebiotic
30 ingredient, due to the reported high resistance of α 1 \rightarrow 2 linkages to the digestive
31 enzymes in humans and animals, as well as to its potential selective stimulation of
32 beneficial bacteria in the large intestine mainly attributed to the two linked glucose units
33 located at the reducing end that reflects the disaccharide kojibiose (2- α -D-
34 glucofuranosyl-D-glucose). These findings could contribute to broaden the use of
35 important agricultural raw materials, such as sucrose or cheese whey permeates, as
36 renewable substrates for enzymatic synthesis of oligosaccharides of nutritional interest.

37

38 **Keywords:** *cheese whey permeate, Leuconostoc mesenteroides B-512F dextransucrase,*
39 *glucosyl-lactose, transglucosylation, kojioligosaccharides.*

40 Introduction

41

42 Development of new strategies for the synthesis of oligosaccharides with
43 functional properties is currently in great demand in the food, pharmaceutical, feed and
44 cosmetic industries, since carbohydrates may find immediate applications as stabilizers,
45 bulking compounds, immunomodulating agents, or prebiotic compounds (1). In this
46 context, the use of enzyme catalysts is a preferred approach to multi-step chemical
47 synthesis considering the high stereo- and regio-selectivity of enzymes (2). Likewise,
48 there is an increasing interest in alternative uses of agricultural and livestock materials,
49 such as sucrose or whey permeates (WPs), as renewable substrates for enzymatic
50 synthesis of valuable saccharides (3, 4). Thus, sucrose is an abundant and low-cost
51 substrate with a great potential for its use as a sustainable raw material for the
52 development of new food ingredients (5) and, particularly, for the synthesis of prebiotic
53 oligosaccharides using glycosyltransferases (6).

54 In addition, WP is an important by-product of the dairy industry containing
55 mainly lactose and salts, which is obtained when cheese or casein whey are subjected to
56 a process of ultrafiltration membrane to concentrate whey proteins. Unlike whey
57 proteins that find immediate food applications, the WP has so far been of little value (7),
58 being its profitable use one of the biggest dairy industry challenge ahead (8). One of the
59 most important uses of whey permeate is the synthesis of galactooligosaccharides
60 (GOS) from transgalactosylation of lactose catalyzed by glycosidases (EC 3.2), being β -
61 galactosidases (EC 3.2.1.23) of microbial origin the most frequently used (4, 9-11). In
62 this sense, it is remarkable to mention that notable differences, in terms of yield and
63 oligosaccharide composition, have been reported between the production of GOS from
64 model systems consisting of lactose in buffered solutions and from WPs. These

65 dissimilarities can be attributed to the influence of other permeate ingredients, such as
66 mineral salts, on the enzymatic reaction (12-15).

67 Despite the broad specificity of glycosidases, their application is often limited by
68 low yields and poor region-selectivity (16). Glycosyltransferases (EC 2.4) are
69 considered as an interesting alternative to glycosidases because they include enzymes
70 capable of transferring glycosyl groups, after hydrolysis of the donor, to water or other
71 acceptors catalyzing, thus, oligosaccharide synthesis (2). Glycosyltransferases are
72 further subdivided, according to the nature of the sugar residue being transferred, into
73 hexosyltransferases (EC 2.4.1), pentosyltransferases (EC 2.4.2), and those transferring
74 other glycosyl groups (EC 2.4.99) (16). Within the subgroup 2.4.1 and under the term
75 glycosyltransferases are considered glucansucrases or fructansucrases. This type of enzymes
76 maintain the regiospecificity and catalyze effective synthesis of various
77 oligosaccharides of different structural nature by using sucrose as glucosyl or fructosyl
78 moiety donor and carbohydrates with low molecular weight acting as acceptors (17, 18).

79 Concretely, dextransucrase (EC 2.4.1.5) is a glucansucrase produced by various
80 species of *Leuconostoc*, *Lactobacillus*, and *Streptococcus* which catalyzes the synthesis
81 of dextran from sucrose and also the transfer of glucose from sucrose (donor) to other
82 carbohydrates (acceptor) by linking mainly an α -(1 \rightarrow 6)-glucosyl bond
83 (transglycosylation reaction) (19, 20). Maltose is recognized as the best acceptor
84 providing the synthesis of a series of potential bioactive oligosaccharides acceptor-
85 products such as panose (6- α -D-glucopyranosylmaltose) and other
86 isomaltooligosaccharides (21, 22). When lactose is the acceptor, only one acceptor
87 product has been reported to be formed, 2- α -D-glucopyranosyl-lactose (22-24).
88 According to its structure, this trisaccharide is an excellent candidate for a new prebiotic
89 ingredient, due to the high resistance of α -(1 \rightarrow 2) linkages to the digestive enzymes in

90 human beings and animals and their selective stimulation of bacteria that are beneficial
91 to the large intestine (25, 26). Nevertheless, to the best of our knowledge, there are no
92 data concerning the production of oligosaccharides from WPs and sucrose by
93 glycosyltransferases.

94 Firstly, this study will address the production and structural characterization by
95 NMR of potentially bioactive oligosaccharides derived from sucrose and lactose
96 catalyzed by dextransucrase of *Leuconostoc mesenteroides* B-512F, to be then applied
97 to the production of oligosaccharides from industrial cheese WPs.

98

99 Materials and methods

100

101 Chemical and reagents

102

103 Dextransucrase from *Leuconostoc mesenteroides* B-512F was purchased from
104 CRITT Bio-Industries (Toulouse, France). Specific activity was 0.4 U mg⁻¹, where 1
105 unit is the amount of enzyme required to perform the transfer of 1 μmol of glucose per
106 minute at a working temperature of 30 °C, a sucrose concentration of 100 g L⁻¹ at pH
107 5.2 with 10 mg L⁻¹ of CaCl₂ · 2H₂O. Sucrose was purchased from Panreac (Barcelona,
108 Spain), and fructose, glucose, raffinose, lactose and leucrose were from Sigma–Aldrich
109 (Steinheim, Germany). Acetonitrile (HPLC grade) was obtained from Lab-scan
110 (Gliwice, Poland). Ultra-pure water quality (18.2 MΩcm) with 1–5 ppb total organic
111 carbon (TOC) and <0.001 EU mL⁻¹ pyrogen levels was produced in-house using a
112 laboratory water purification Milli-Q Synthesis A10 system from Millipore (Billerica,
113 MA). All other chemicals were of analytical grade.

114

115 Physical-chemical characterization of cheese whey permeates

116

117 Two different industrial bovine cheese whey permeate powders (WPs) were
118 kindly supplied by the dairy industries García Baquero (Alcázar de San Juan, Spain)
119 and Reny Picot (Navia, Spain).

120 The pH of both WPs was measured using a pH meter (MP 230, Mettler-Toledo,
121 Barcelona, Spain) at a concentration of 50 mg mL⁻¹.

122 The dry matter (DM) content of WPs was gravimetrically determined by drying
123 the samples in a conventional oven at 102 °C until constant weight, according to the
124 AOAC method (27).

125 Ion composition of the WPs was determined using an ICP-MS ELAN 6000
126 Perkin Elmer Sciex instrument at the Servicio Interdepartamental de Investigación
127 (SIdI-UAM) of Madrid. Either a semiquantitative analysis or a quantitative analysis of
128 the elements of interest using the external calibration method and internal standards to
129 correct instrumental drift were carried out (28).

130 The total protein content was determined in WPs aqueous dissolutions (300 mg
131 mL⁻¹) according to Bradford's dye binding method, using bovine serum albumin (BSA)
132 as standard (29).

133

134 Oligosaccharide synthesis

135

136 Oligosaccharide synthesis in the presence of sucrose (donor) and lactose
137 (acceptor) was carried out by incubating 1 or 2 mg (0.4 or 0.8 U, respectively) of
138 dextransucrase per mL at 30 °C in 20 mM sodium acetate buffer with 0.34 mM CaCl₂ at
139 pH 5.2. Product formation was investigated by taking aliquots from the reaction mixture

140 at suitable time intervals up to 48 h. The enzyme was inactivated by heating at 100 °C
141 for 5 min and inactivated samples were then diluted with acetonitrile:water (50:50, v:v),
142 filtered using a 0.45 µm syringe filter (Symta, Madrid, Spain), and analyzed by two
143 different LC systems as described below. In order to investigate the influence of
144 synthetic conditions on the formation of the oligosaccharide of interest, the reactions
145 were done at two different concentrations of substrates, i.e. sucrose:lactose 45:20 and
146 30:30, expressed in g/100 mL, leading to two different molar ratios of donor/acceptor
147 (2.25:1 and 1:1).

148 Once factors such as enzyme charge and sucrose:lactose ratio were studied, the
149 enzymatic reactions were performed with sucrose:cheese WPs mixtures at 30 and 40 °C.
150 Considering the lactose content measured in both WPs, enzymatic reactions were
151 carried out at sucrose:lactose 30%:30%, in 20mM sodium acetate buffer (pH 5.2) for 48
152 hours. When it was necessary, pH was adjusted to 5.2 with acetic acid.

153 Moreover, the effect of the major ions present in the WPs on the *L.*
154 *mesenteroides* B-512F dextransucrase activity was evaluated. Thus, 8 µl of 1M
155 Na₂HPO₄ and/or 20 µl of 1M K₂HPO₅ (both solutions with a pH value adjusted to 5.2)
156 were added to 1 mL of sucrose:lactose 30%:30% previously dissolved in 20mM sodium
157 acetate buffer (pH 5.2). The enzymatic reactions were carried out as described above at
158 30 °C for 48 hours.

159

160 Chromatographic determination of carbohydrates

161

162 *Liquid chromatography with refractive index detector*

163

164 The synthesized oligosaccharides were analysed by liquid chromatography with
165 refractive index detector (HPLC-RID) on an Agilent Technologies 1220 Infinity LC
166 System – 1260 RID (Boeblingen, Germany). The separation of carbohydrates was
167 carried out with a Kromasil® (100-NH₂) column (250 x 4.6 mm, 5 µm particle size)
168 (Akzo Nobel, Brewster, NY) using acetonitrile:water (75:25, v:v) as the mobile phase
169 and eluted in isocratic mode at a flow rate of 1.0 mL min⁻¹ for 50 min. Injection volume
170 was 50 µL (800 µg of total carbohydrates). Data acquisition and processing were
171 performed using the Agilent ChemStation software (Agilent Technologies, Boeblingen,
172 Germany).

173 Carbohydrates in the reaction mixtures were initially identified by comparing the
174 retention times (t_R) with those of standard sugars. Quantitative analysis was performed
175 by the external standard method, using calibration curves in the range 0.01-10 mg for
176 fructose (quantification of monosaccharides), sucrose, lactose and leucrose
177 (disaccharides) and raffinose (quantification of trisaccharides). All analyses were
178 carried out in triplicate. Determination coefficients obtained from these calibration
179 curves, which were linear over the range studied, were high ($R^2 > 0.999$).
180 Reproducibility of the method was estimated on the basis of the intra-day and inter-day
181 precision, calculated as the relative standard deviation (*RSD*) of concentrations of
182 oligosaccharide standards obtained in $n \geq 5$ independent measurements, obtaining *RSD*
183 values below 10% in all cases.

184

185 *High performance anion exchange chromatography with pulsed amperometric*
186 *detection*

187 In order to increase sensitivity, some of the samples were also analyzed by high
188 performance anion exchange chromatography with pulsed amperometric detection

189 (HPAEC-PAD) on an IC 2500 Dionex System consisting of a GP50 gradient pump and
190 an ED50 electrochemical detector with a gold working electrode and Ag/AgCl reference
191 electrode. Data acquisition and processing were performed with a Chromeleon version
192 6.7 software (Dionex Corp., Sunnyvale, CA). For eluents preparation, MilliQ water,
193 50% (w:v) NaOH, and NaOAc (Panreac, Barcelona, Spain) were used. All eluents were
194 degassed by flushing with helium for 25 min. Separations were performed following the
195 method described by Spletchna et al. (30). Elution was at room temperature on a
196 CarboPac PA-1 column (250 × 4 mm) connected to a CarboPac PA-1 (50 × 4 mm)
197 guard column. The elution, at a flow rate of 1 mL min⁻¹, was in gradient using a
198 combination of three eluents: A (100 mM NaOH), B (100 mM NaOH and 50 mM
199 NaOAc), and C (100 mM NaOH and 1 M NaOAc). The gradient used was 100% A
200 from 0 to 20 min and 0-100% B from 20 to 70 min. After each run, the column was
201 washed for 10 min with 100% C and re-equilibrated for 15 min with the starting
202 conditions of the employed gradient.

203

204 *Gas chromatography with mass spectrometry detection*

205

206 Selected samples were also analyzed by gas chromatography with a mass
207 spectrometry detector (GC-MS) on an Agilent Technologies 7890A gas chromatograph
208 coupled to a 5975C MSD quadrupole mass detector (Agilent Technologies,
209 Wilmington, DE). The trimethylsilyloximes (TMSO), prepared as described by Sanz et
210 al. (31), were separated using an HP-5 MS fused-silica capillary column (30 m × 0.25
211 mm i.d. × 0.25 μm film thickness) coated with 5% phenylmethylsilicone (J&W
212 Scientific, CA). The helium flow rate was 1 mL min⁻¹. The initial oven temperature was
213 180 °C and increased to 315 °C at a heating rate of 3 °C min⁻¹ and held for 20 min. The

214 injector temperature was 280 °C and injections were made in the split mode (1:40).
215 Mass spectrometer was operated in electronic impact mode at 70 eV. Mass spectra were
216 acquired using Agilent ChemStation MSD software (Wilmington, DE).

217 Identification of TMSO derivatives of carbohydrates was carried out by
218 comparison of their retention indices and mass spectra with those of standard
219 compounds previously derivatized.

220

221 Purification and structural characterization of the main acceptor product by Nuclear
222 Magnetic Resonance

223

224 Considering the absence of commercially available standard for the main
225 synthesised oligosaccharide, this trisaccharide was isolated and purified by HPLC-RID
226 from sucrose:lactose mixtures after 24 hours of enzymatic reaction and using a semi-
227 preparative column Kromasil[®] (100-NH₂) column (250 × 10 mm, 5 µm particle size)
228 (Akzo Nobel, Brewster, NY). Thus, 500 µL of reaction mixtures (15 mg of total
229 carbohydrates) was repeatedly eluted with acetonitrile:water (75:25, v:v) as the mobile
230 phase at a flow rate of 5 mL min⁻¹, and fractions corresponding to the main synthesised
231 oligosaccharide were manually collected, pooled, and evaporated in a rotatory
232 evaporator R-200 (Büchi, Switzerland) below 25 °C for its subsequent characterization.

233 Structure elucidation of the purified oligosaccharide was accomplished by
234 Nuclear Magnetic Resonance spectroscopy (NMR). NMR spectra were recorded at 298
235 and 313 K, using D₂O as the solvent, on a Varian SYSTEM 500 NMR spectrometer (¹H
236 500 MHz, ¹³C 125 MHz) equipped with a 5 mm HCN cold probe. Chemical shifts of ¹H
237 (δ_H) and ¹³C (δ_C) in ppm were determined relative to an external standard of sodium [2,
238 2, 3, 3-²H₄]-3-(trimethylsilyl)-propanoate in D₂O (δ_H 0.00 ppm) and 1, 4-dioxane (δ_C

239 67.40 ppm) in D₂O, respectively. One-dimensional NMR experiments (¹H, and ¹³C)
240 were performed using standard Varian pulse sequences. Two-dimensional [¹H, ¹H]
241 NMR experiments (gCOSY and TOCSY) were carried out with the following
242 parameters: a delay time of 1 s, a spectral width of 1675.6 Hz in both dimensions, 4096
243 complex points in t₂ and 4 transients for each of 128 time increments, and linear
244 prediction to 256. The data were zero-filled to 4096 × 4096 real points. Two-
245 dimensional [¹H-¹³C] NMR experiments (gHSQC and gHMBC) used the same ¹H
246 spectral window, a ¹³C spectral windows of 30165 Hz, 1 s of relaxation delay, 1024 data
247 points, and 128 time increments, with a linear prediction to 256. The data were zero-
248 filled to 4096 × 4096 real points. Typical numbers of transients per increment were 4
249 and 16, respectively.

250

251 **Results and Discussion**

252

253 Optimization of enzymatic synthesis conditions by using model systems based on 254 sucrose:lactose mixtures

255

256 Given that the optimum pH (5.2) and temperature (30 °C) of *Leuconostoc*
257 *mesenteroides* B-512F dextransucrase for glucansucrase activity have been previously
258 well established (32, 33), the production of oligosaccharides was studied as a function
259 of the ratio of sucrose to lactose, and the concentration of enzyme as described below.

260 A sucrose concentration of 45% and a lactose concentration of 20%, expressed
261 in g/100 mL, were initially employed. These values were based on: i) the higher the
262 concentration of substrates, the greater the inhibition of formation of dextran and the
263 greater the formation of acceptor-reaction products (20), ii) previous studies described

264 an initial sucrose concentration of 45% to be optimal for glucooligosaccharides
265 production from sucrose:maltose mixtures (25). Furthermore, taking into account that
266 an increase in the ratio of acceptor to sucrose leads to a dramatic decrease in the amount
267 of dextran with a concomitant increase in the amount of acceptor products (20),
268 enzymatic reactions with 30% of sucrose and 30% of lactose, expressed in g/100 mL,
269 were also evaluated. Finally, both sets of samples were treated with two different
270 concentrations of dextransucrase (0.4 and 0.8 U mL⁻¹) under standard reaction condition
271 (at pH 5.2 and 30°C for 48 h). Results obtained with 0.4 U mL⁻¹ of enzyme showed that
272 sucrose was not totally consumed after 48 hours of reaction and lower levels of
273 oligosaccharides were formed (data not shown) as compared to the equivalent reaction
274 mixtures treated with 0.8 U mL⁻¹ of dextransucrase. These results are in good agreement
275 with data reported by Robyt (20) who indicated that keeping a constant 1:1 ratio of
276 maltose to sucrose, an increase in the concentration of enzyme gave rise to a decrease in
277 dextran and to an increase in the amount of acceptor-reaction products. Thus, all results
278 shown in this work correspond to a concentration of enzyme of 0.8 U mL⁻¹.

279 HPLC-RID chromatograms of the enzymatic reactions mixtures at 0 and 24
280 hours with sucrose:lactose mixtures at concentrations of 45%:20% and 30%:30% are
281 shown in **Figures 1A and 1B**. As expected, at the initial time only two well-resolved
282 peaks corresponding to sucrose (peak 3, t_R 9.7 min) and lactose (peak 5, t_R 12.5 min)
283 were detected. Nevertheless, after 24 hours of reaction four new peaks clearly appeared
284 (peaks 1, 2, 4 and 7), whilst the peak corresponding to lactose substantially decreased,
285 which is indicative of its capacity as acceptor. Likewise, at sucrose:lactose mixtures of
286 30%:30% the peak corresponding to sucrose was detected at trace levels, showing that
287 sucrose was readily consumed after 24 hours of reaction by the hydrolytic action of the
288 dextransucrase (**Figure 1B**). At 24 hours of reaction, the less retained peaks (1 and 2)

289 corresponded to fructose and glucose (t_R 6.9 and 7.8 min, respectively), and they were
290 derived from the hydrolysis of sucrose. Furthermore, fructose was much more abundant
291 than glucose, indicating that glucose was efficiently transferred. The other two new
292 peaks (named 4 and 7, t_R 10.7 and 20.3 min) were detected in the eluting area of di- and
293 trisaccharides, respectively, suggesting that both of them were acceptor-reaction
294 products. Finally, a minor set of peaks (termed 6) was also detected around 16 min, that
295 might correspond to minor acceptor-reaction products (**Figure 1**).

296 Prior to quantification, peak 4 corresponding to a disaccharide (t_R 10.7 min,
297 **Figure 1**) was identified as leucrose (5-*O*- α -D-glucopyranosyl-D-fructopyranose) by
298 GC-MS analysis of its corresponding trimethylsilyloxime (TMSO) and comparison with
299 a standard previously derivatized (spectrum not shown). The mass spectrum of leucrose
300 was characterized by the m/z ions (by decreasing order of abundance): 361, 204, 217,
301 538, 271 and 243. The formation of leucrose proves that fructose can also act as
302 acceptor in the dextransucrase-catalyzed reactions (16). Likewise, the major acceptor-
303 reaction product eluting on the trisaccharide area (peak 7, t_R 20.3, **Figure 1**) could
304 correspond to glucosyl-lactose as it has been previously reported in studies on the
305 enzymatic synthesis of oligosaccharides in a lactose-sucrose medium either inoculated
306 with *Betacoccus arabinosaceus* cultures (23, 24) or after addition of dextransucrase
307 from *L. mesenteroides* B-512F (22). In order to confirm this fact, the purification and
308 exhaustive characterization by NMR of this compound will be discussed in the next
309 section.

310 **Figures 2A** and **2B** illustrate the concentration of sucrose, lactose, leucrose and
311 glucosyl-lactose upon the enzymatic reaction time at sucrose:lactose ratios 45%:20%
312 and 30%:30%, respectively. The maximum formation of the main acceptor-reaction
313 product was achieved at 24-32 hours and then remained practically constant to the end

314 of the enzymatic reaction. This increase and subsequent plateau coincided with the
315 gradual decrease of lactose observed up to 24 hours of reaction (loss of 40-49% of
316 lactose in weight respect to the initial amount) and the posterior plateau, confirming that
317 lactose was the acceptor molecule for the formation of the main trisaccharide (**Figures**
318 **2A** and **2B**). By comparing both set of reaction mixtures, it can be inferred that whilst
319 the maximum amount of formed glucosyl-lactose was fairly similar in both cases (120-
320 130 mg mL⁻¹), the formation of leucrose (which could be considered as an unwanted by-
321 product) was two-fold lower in the presence of 30% of sucrose and 30% of lactose than
322 when 45% sucrose and 20% lactose mixture was used (i.e., 27 and 54 mg mL⁻¹ of
323 leucrose, respectively, after 32 hours of reaction). This result can be explained by the
324 fact that the formation of leucrose by action of dextransucrase is favoured at high
325 fructose concentration (16). This means that the higher the concentration of sucrose, the
326 higher the concentration of fructose in the reaction medium and, consequently, the
327 greater the formation of leucrose.

328 Overall, the maximum yields of glucosyl-lactose and leucrose were 23.3% and
329 5.1%, based on total carbohydrates, respectively, when 30% of sucrose and 30% of
330 lactose were initially present in the enzymatic reaction. This means that a yield of
331 approximately 47% of glucosyl-lactose (in weight respect to the initial amount of
332 lactose) was attained, being this value markedly higher than those previously reported
333 (22, 23). This can be attributed to the fact that in the present study the initial substrates
334 (sucrose and lactose) were much more concentrated than those employed by those
335 authors.

336 Considering that electrochemical detectors such as pulsed-amperometric detector
337 (PAD) have a higher sensitivity than the refractive index detector (RID), samples taken
338 after 24 hours of enzymatic reaction in the presence of sucrose:lactose mixtures at

339 30%:30% and 45%:20% were analysed by HPAEC-PAD in order to find other minor
340 oligosaccharides not detected by HPLC-RID. In addition to the compounds previously
341 detected by HPLC-RID, HPAEC-PAD chromatograms showed a series of peaks eluting
342 from 28 to 68 min which could correspond to other minor acceptor-reaction products of
343 degree of polymerization equal to or above 3 (**Figure 3**). The formation of minor
344 acceptor-reaction products explained that the loss of moles of lactose was slightly
345 higher than the production of moles of the main synthesized oligosaccharide (glucosyl-
346 lactose). As an example, at 24 hours of reaction 0.32 mol L⁻¹ of lactose were lost
347 whereas 0.24 mol L⁻¹ of glucosyl-lactose were produced.

348

349 *NMR characterization of glucosyl-lactose*

350

351 In order to elucidate the complete structure of the main synthesized
352 oligosaccharide, the enzymatic reaction mixture at 24 hours (sucrose:lactose, 30%:30%)
353 was repeatedly separated by HPLC-RID using a semi-preparative column and the
354 corresponding chromatographic peak was manually collected to be further analysed by
355 NMR. Then, unequivocal structural elucidation of this compound was carried out by the
356 combined use of 1D and 2D [¹H, ¹H] and [¹H-¹³C] NMR experiments (gCOSY,
357 TOCSY, multiplicity-edited gHSQC and gHMBC). Experiments were carried out at
358 313K to avoid the superposition of HDO and one of the anomeric protons. ¹H and ¹³C
359 NMR chemical shifts are given in **Table 1**. To the best of our knowledge, this is the first
360 report of ¹H and ¹³C-NMR full assignments for this compound and full set of spectra are
361 collected in Supporting Information.

362 ¹H NMR spectrum of the trisaccharide showed six doublets in the anomeric
363 region. The TOCSY experiment revealed the ¹H signals of four glucose and two

364 galactose residues (**Figure 4**). These results are compatible with an anomeric mixture at
365 one of glucose residues. In addition, ^{13}C NMR spectrum displayed two sets of 18
366 resonances. The major set of resonances, corresponding to the most populated isomer,
367 contained three anomeric carbons at δ 103.99, 98.81 and 97.07. The minor set of
368 resonances also contained three anomeric carbons at δ 103.90, 97.46 and 90.23. A
369 multiplicity-edited gHSQC spectrum was used to link the carbon signals to the
370 corresponding proton resonances. So, taking the anomeric carbons as starting point, one
371 residue of β -galactose (Gal, $J(\text{H1},\text{H2}) = 6.3$ Hz), one residue of β -glucose (Glu,
372 $J(\text{H1},\text{H2}) = 8.0$ Hz), and one residue of α -glucose (Glu', $J(\text{H1},\text{H2}) = 3.9$ Hz) were
373 identified for the major isomer and one residue of β -galactose (Gal, $J(\text{H1},\text{H2}) = 6.1$
374 Hz), one residue of α -glucose (Glu, $J(\text{H1},\text{H2}) = 3.5$ Hz) and one residue of α -glucose
375 (Glu', $J(\text{H1},\text{H2}) = 3.8$ Hz) for the minor isomer.

376 The position of glycosidic linkages was analyzed as follows. For the major
377 isomer, gHMBC showed correlations between the Gal-H1 anomeric proton (4.33 ppm)
378 and the Glu- C4 carbon (79.73 ppm) and between the Glu'-C1 anomeric carbon (98.81
379 ppm) and the Glu- H2 proton (3.31 ppm). For the minor-isomer, gHMBC showed
380 correlations between the Gal-H1 anomeric proton (4.35 ppm) and the Glu- C4 carbon
381 (79.43 ppm) and between the Glu'-C1 anomeric carbon (97.46 ppm) and the Glu- H2
382 proton (3.58 ppm). So, the major compound was identified as *O*- β -D-galactopyranosyl-
383 (1 \rightarrow 4)-*O*-[α -D-glucopyranosyl-(1 \rightarrow 2)]- β -D-glucopyranose and the minor isomer as *O*-
384 β -D-galactopyranosyl-(1 \rightarrow 4)-*O*-[α -D-glucopyranosyl-(1 \rightarrow 2)]- α -D-glucopyranose. The
385 anomeric mixture was 58% (β -anomer) and 42% (α -anomer) by integration of the
386 anomeric protons. Both anomers can be defined as 2- α -D-glucopyranosyl-lactose in
387 order to simplify the trisaccharide nomenclature.

388

389 *Production of 2- α -D-glucoopyranosyl-lactose from cheese whey permeates*

390

391 In order to study the enzymatic synthesis of 2- α -D-glucoopyranosyl-lactose from
392 WPs, two different industrial cheese WPs (WP1 and WP2) were used. A preliminary
393 physical-chemical characterization of both permeates was carried out in order to explain
394 possible dissimilarities in their behaviour to the enzymatic synthesis (**Table 2**). Both
395 cheese WPs had lactose contents within the range established for typical composition of
396 whey permeate (34), whilst the main identified metal ions were those derived from the
397 most abundant mineral salts detected in whey permeates such as calcium, magnesium,
398 sodium, potassium and phosphorus. Nevertheless, there were notable differences in the
399 total mineral content which were principally attributed to the significantly high levels of
400 sodium and, especially, phosphorus and potassium present in WP2 (**Table 2**). Lastly,
401 permeates also contained trace minerals including boron, zinc, bromine, rubidium,
402 strontium, iodine, titanium and barium. In addition, differences in the pH values
403 measured for both WPs could suggest that different cheese technological processes were
404 employed.

405 For the enzymatic synthesis of 2- α -D-glucoopyranosyl-lactose, sucrose was
406 added in both whey permeates as donor at a 1:1 weight ratio considering the
407 concentration of lactose previously determined (**Table 2**). Thus, the enzymatic reaction
408 started with 30% of sucrose and 30% of lactose, since these conditions gave the best
409 result for higher production of the oligosaccharide of interest with the model systems.
410 Likewise, the pH of both whey permeates was adjusted to a value of 5.2 (optimum pH
411 for the activity of dextransucrase).

412 HPLC-RID profiles of enzymatic reactions with both WPs were similar to those
413 obtained for the model systems (sucrose:lactose, 30%:30%). Thus, the main detected

414 carbohydrates after 24 hours of enzymatic reaction corresponded to (by decreasing
415 order of abundance): lactose, 2- α -D-glucopyranosyl-lactose, fructose, leucrose and
416 glucose (**Figure 1C**). In addition, as it occurred for the model systems, a series of peaks
417 at trace levels, probably corresponding to minor acceptor-reaction products (peak 6),
418 were detected at 16-17 min.

419 **Figures 2C and 2D** show the concentration of sucrose, lactose, leucrose and 2-
420 α -D-glucopyranosyl-lactose upon the enzymatic reaction time with both type of cheese
421 WPs. Regarding leucrose, maximum yields were 4.5% at 24 hours, and 5.3% at 32
422 hours for WP1 and WP2, respectively (in weight respect to total carbohydrates). In good
423 agreement with the results obtained for the model systems, the maximum formation of
424 2- α -D-glucopyranosyl-lactose in both cheese WPs was attained at 24 hours of reaction,
425 and then it remained constant to the end of the reaction whilst lactose content also
426 exhibited a gradual decrease for the first 24 hours of reaction. However, when both WPs
427 were compared, WP2 presented higher levels of 2- α -D-glucopyranosyl-lactose than
428 WP1. Thus, the highest yields of 2- α -D-glucopyranosyl-lactose were 42% (122 mg mL⁻¹)
429 ¹) and 52.4% (155 mg mL⁻¹) for WP1 and WP2, respectively (in weight respect to the
430 initial amount of lactose). These dissimilarities in yield of oligosaccharide synthesis
431 could be partially attributed to the substantial differences found in the mineral salts
432 content for both assayed WP, where WP2 had higher levels of sodium, phosphorus and
433 potassium than WP1 (**Table 2**). Although diverse studies have shown that the effect of
434 metal ions on oligosaccharide synthesis largely depends on the source from which the
435 enzyme has been isolated (12) and on the substrate used in the assay (35), cations, such
436 as sodium, potassium and magnesium, have been described to modify the
437 transgalactosylation activity of a β -galactosidase from bacterial origin (36). Moreover,

438 the yield of 2- α -D-glucopyranosyl-lactose produced from WP2 was also higher than
439 that previously obtained from the sucrose:lactose (30%:30%) model system.

440 Taking into account the above results, the effect of the major ions, that were
441 present in higher amounts in the WP2 than in the WP1 (i.e., sodium, potassium and
442 phosphorus, **Table 2**), on the *L. mesenteroides* B-512F dextransucrase activity was
443 evaluated. These studies were carried out using sucrose and lactose mixtures
444 (30%:30%) and adding the major ions at the equivalent concentration as they were
445 present in the WP2. Results showed that either the combined or the unique presence of
446 sodium, phosphorous and/or potassium increased the sucrose hydrolysis rate but the
447 content of 2- α -D-glucopyranosyl-lactose was not significantly modified (data not
448 shown). In this sense, it should not be ruled out that the trace minerals contained in the
449 studied WPs (**Table 2**) might also have an effect on the transfer rate of the
450 dextransucrase and, therefore, on the content of 2- α -D-glucopyranosyl-lactose.

451 Furthermore, the enzymatic reaction was also performed at 40 °C in addition to
452 30 °C in order to attempt a reduction in the reaction time. However, at 40 °C, there were
453 not significant increases in yield of 2- α -D-glucopyranosyl-lactose nor the reaction time
454 was shortened (data not shown), probably due to 40 °C exceeds the optimum
455 temperature range of dextransucrase from *L. mesenteroides* (32, 33).

456

457 Given that colonic microbiota has an extraordinary contribution to well-being
458 and health of the host, there is a growing interest in identifying functional dietary
459 compounds capable of modulating the metabolic activities and composition of the
460 intestinal microbiota (37). An alternative to probiotics is the use of prebiotics, which
461 have been defined as “non-digestible ingredients that are selectively fermented and
462 allows specific changes, both in the composition and/or activity in the gastrointestinal

463 microflora that confers benefits upon host wellbeing and health” (38). The trisaccharide,
464 2- α -D-glucopyranosyl-lactose, synthesised and characterized in this work is an
465 excellent candidate for a new prebiotic ingredient since, on the one hand, it has been
466 shown the high resistance of α -(1 \rightarrow 2) linkages to *in vitro* and *in vivo* gastrointestinal
467 digestion (25, 39, 40). On the other hand, this trisaccharide is an oligosaccharide
468 derived from kojibiose (2-*O*- α -D-glucopyranosyl-D-glucose) as it contains this
469 disaccharide at the reducing end of the structure (**Table 1** and **Figure 4**). Thus, previous
470 studies demonstrated the growth of 6 strains belonging to *Bifidobacterium*,
471 *Lactobacillus* and *Eubacterium* during the incubation of kojibiose with pure cultures.
472 Likewise, kojioligosaccharides with a degree of polymerization of 3 and 4 were utilized
473 by fewer strains but more selectively than kojibiose (40). In this context, Sanz et al. (41)
474 studied the influence of glycosidic linkages and monosaccharide composition on the
475 selectivity of microbial fermentation in a wide range of disaccharides following their *in*
476 *vitro* fermentation by using human fecal batch cultures; strikingly, kojibiose showed the
477 greatest prebiotic index of the 20 assayed disaccharides, identifying this disaccharide
478 and its potential derived-oligosaccharides as one of the most promising novel
479 carbohydrates that could be beneficial to gut health.

480

481 To conclude, in this study, it has been shown the capacity of dextransucrase (EC
482 2.4.1.5) from *Leuconostoc mesenteroides* B-512F to catalyze the high-yield synthesis of
483 potentially bioactive 2- α -D-glucopyranosyl-lactose, characterized by NMR, in the
484 presence of sucrose and cheese whey permeates. These findings could contribute to
485 broaden the use of these important agricultural raw materials as renewable substrates for
486 enzymatic synthesis of oligosaccharides of nutritional interest.

487

488 *Acknowledgements*

489

490 This work has been financed by projects POII10-0178-4685 from Junta de
491 Comunidades de Castilla-La Mancha and European regional development fund (ERDF),
492 and projects AGL2011-27884 and Consolider Ingenio 2010 FUN-C-FOOD CSD2007-
493 00063 both from Ministerio de Ciencia e Innovación. M. Díez-Municio thanks CSIC for
494 a JAE predoctoral fellowship. We thank Francisco José Redondo and Ricardo González
495 from García Baquero and Reny Picot, respectively, for kindly providing us with cheese
496 whey permeates. We also thank Inmaculada Rivas and Zaida Gil of the ICP-MS service
497 of SIDI-UAM for help in the physical-chemical characterization of cheese whey
498 permeates.

499 Literature cited

- 500 1. André, I.; Potocki-Véronèse, G.; Morel, S.; Monsan, P.; Remaud-Siméon, M.
501 Sucrose-utilizing transglucosidases for biocatalysis. *Top. Curr. Chem.* **2010**, 294, 25–
502 48.
- 503 2. Monsan, P.; Paul, F. Enzymatic synthesis of oligosaccharides. *FEMS Microbiol. Rev.*
504 **1995**, 16, 187-192.
- 505 3. Baciú, I. E. Extracted sugar-beet pulp and sucrose, two renewable materials as “hot”
506 substrates for enzymatic synthesis of valuable saccharides. PhD Dissertation;
507 *Technical University Braunschweig*. **2005**.
- 508 4. Adamczak, M.; Charubin, D.; Bednarski, W. Influence of reaction medium
509 composition on enzymatic synthesis of galactooligosaccharides and lactulose from
510 lactose concentrates prepared from whey permeate. *Chemical Papers*. **2009**, 63, 111-
511 116.
- 512 5. Khan, R. Sucrose: Its potential as a raw material for food ingredients and for
513 chemicals. In *Sucrose; properties and applications*. Mathlouthi, M., Reiser, P., Eds.;
514 Blackie A&P: London, U.K., 1995; pp. 264-278.
- 515 6. Monsan, P.F.; Ouarné, F. Oligosaccharides derived from sucrose. In *Prebiotics and*
516 *probiotics science and technology*. Charalampopoulos, D., Rastall, R.A, Eds.; Springer-
517 Verlag: New York, NY, 2009; Vol.1, pp. 293-336.
- 518 7. Barile, D.; Tao, N.; Lebrilla, C.B.; Coisson, J.-D.; Arlorio, M.; German, J.B.
519 Permeate from cheese whey ultrafiltration is a source of milk oligosaccharides. *Int.*
520 *Dairy J.* **2009**, 19, 524-530.
- 521 8. 3A Business Consulting. *Global Opportunities for Whey and Lactose Ingredients*
522 *2010-2014*. 2010. Available from: <http://www.3abc.dk/Report%20information%20->

- 523 %20Global%20Opportunities%20for%20Whey%20and%20Lactose%20Ingredients%20
524 02010-2014.pdf
- 525 9. Rustom, I.Y.S.; Foda, M.I.; Lopez-Leiva, M. Formation of oligosaccharides from
526 whey UF-permeate by enzymatic hydrolysis: analysis of factors. *Food Chem.* **1998**, *62*,
527 141-147.
- 528 10. Goulas, A.; Tzortzis, G.; Gibson, G.R. Development of a process for the production
529 and purification of α - and β -galactooligosaccharides from *Bifidobacterium bifidum*
530 NCIMB 41171. *Int. Dairy J.* **2007**, *17*, 648-656.
- 531 11. Splechtna, B.; Nguyen, T.H.; Haltrich, D. Comparison between discontinuous and
532 continuous lactose conversion processes for the production of prebiotic galacto-
533 oligosaccharides using β -galactosidase from *Lactobacillus reuteri*. *J. Agric. Food*
534 *Chem.* **2007**, *55*, 6772-6777.
- 535 12. Mozaffar, Z.; Nakanishi, K.; Matsuno, R. Formation of oligosaccharides during
536 hydrolysis of lactose in milk using β -Galactosidase from *Bacillus circulans*. *J. Food Sci.*
537 **1985**, *50*, 1602-1606.
- 538 13. Hellerová, K.; Čurda, L. Influence of type of substrate and enzyme concentration on
539 formation of galacto-oligosaccharides. *Czech J. Food Sci.* **2009**, *27*, 372-374.
- 540 14. Gosling, A.; Alftrén, J.; Stevens, G.W.; Barber, A.R.; Kentish, S.E.; Gras, S.L.
541 Facile pretreatment of *Bacillus circulans* β -galactosidase increases the yield of
542 galactosyl oligosaccharides in milk and lactose reaction systems. *J. Agric. Food Chem.*
543 **2010**, *57*, 11570-11574.
- 544 15. Povedicová, K.; Curda, L.; Misún, D.; Dryáková, A.; Diblíková, L. Preparation of
545 galacto-oligosaccharides using membrane reactor. *J. Food Eng.* **2010**, *99*, 479-484.

- 546 16. Plou, F.J.; Martín, M.T.; de Segura, A.G.; Alcalde, M.; Ballesteros, A.
547 Glucosyltransferases acting on starch or sucrose for the synthesis of oligosaccharides.
548 *Can. J. Chem.* **2002**, *80*, 743-752.
- 549 17. Monchois, V.; Willemot, R.M.; Monsan, P. Glucansucrases: mechanism of action
550 and structure–function relationships. *FEMS Microbiol. Rev.* **1999**, *23*, 131-151.
- 551 18. Monsan, P.; Remaud-Siméon, M.; André, I. Transglucosidases as efficient tools for
552 oligosaccharide and glucoconjugate synthesis. *Curr. Opin. Microbiol.* **2010**, *13*, 293-
553 300.
- 554 19. Robyt, J.F.; Eklund, S.H. Stereochemistry involved in the mechanism of action of
555 dextransucrase in the synthesis of dextran and the formation of acceptor products.
556 *Bioorg. Chem.* **1982**, *11*, 115-132.
- 557 20. Robyt, J.F. Mechanisms in the glucansucrase synthesis of polysaccharides and
558 oligosaccharides from sucrose. *Adv. Carbohydr. Chem. Biochem.* **1995**, *51*, 133-168.
- 559 21. Robyt, J.F.; Walseth, T.F. The mechanism of acceptor reactions of *Leuconostoc*
560 *mesenteroides* B-512F dextransucrase. *Carbohydr. Res.* **1978**, *61*, 433-445.
- 561 22. Robyt, J.F.; Eklund, S.H. Relative, quantitative effects of acceptors in the reaction
562 of *Leuconostoc mesenteroides* B-512F dextransucrase. *Carbohydr. Res.* **1983**, *121*, 279-
563 286.
- 564 23. Bourne, E.J.; Hartigan, J.; Weigel, H. 469. Mechanism of the enzymic synthesis of a
565 branched trisaccharide containing the α -1:2-glucosidic linkage. *J. Chem. Soc.* **1959**,
566 2332-2337.
- 567 24. Bailey, R.W.; Barker, S.A.; Bourne, E.J.; Stacey, M. Enzymic Synthesis of a
568 'Branched' Trisaccharide. *Nature.* **1955**, *176*, 1164-1165.

- 569 25. Remaud-Siméon, M.; Willemot, R.M.; Sarçabal, P.; Potocki de Montalk, G.;
570 Monsan, P. Glucansucrases: molecular engineering and oligosaccharide synthesis. *J.*
571 *Molec. Catal. B: Enzym.* **2000**, *10*, 117-128.
- 572 26. Sanz, M.L.; Côté, G.L.; Gibson, G.R.; Rastall, R.A. Influence of glycosidic linkages
573 and molecular weight on the fermentation of maltose-based oligosaccharides by human
574 gut bacteria. *J. Agric. Food Chem.* **2006**, *54*, 9779-9784.
- 575 27. Association of Official Analytical Chemists; by Helrich, K. AOAC method 950.01,
576 Official methods of analysis of AOAC; Arlington, VA,: 1990; Vol.1 pp. 684.
- 577 28. Zuluaga, J.; Rodríguez, N.; Rivas-Ramirez, I.; de la Fuente, V.; Rufo, L.; Amils, R.
578 An improved semiquantitative method for elemental analysis of plants using inductive
579 coupled plasma mass spectrometry. *Biol. Trace Elem. Res.* **2011**, 1-16.
- 580 29. Bradford, M.M. A rapid and sensitive method for the quantitation of microgram
581 quantities of protein utilizing the principle of protein-dye binding. *Anal. Biochem.* **1976**,
582 *72*, 248-254.
- 583 30. Splechtna, B.; Nguyen, T.; Steinböck, M.; Kulbe, K.D.; Lorenz, W.; Haltrich, D.
584 Production of prebiotic galacto-oligosaccharides from lactose using β -galactosidases
585 from *Lactobacillus reuteri*. *J. Agric. Food Chem.* **2006**, *54*, 4999-5006.
- 586 31. Sanz, M.L.; Sanz, J.; Martinez-Castro, I. Gas chromatographic-mass spectrometric
587 method for the qualitative and quantitative determination of disaccharides and
588 trisaccharides in honey. *J. Chromatogr., A.* **2004**, *1059*, 143-148.
- 589 32. Kim, D.; Robyt, J.F. Production and selection of mutants of *Leuconostoc*
590 *mesenteroides* constitutive for glucansucrases. *Enzyme Microb. Technol.* **1994**, *16*, 659-
591 664.
- 592 33. Kim, M.; Day, D.F. Optimization of oligosaccharide synthesis from cellobiose by
593 dextransucrase. *Appl. Biochem. Biotechnol.* **2008**, 707-716.

- 594 34. U.S. Dairy Export Council. *Reference Manual for U.S. Whey and Lactose Products*.
595 2004; Available from:
596 http://usdec.files.cmsplus.com/PDFs/2008ReferenceManuals/Whey_Lactose_Reference
597 [_Manual_Complete2_Optimized.pdf](http://usdec.files.cmsplus.com/PDFs/2008ReferenceManuals/Whey_Lactose_Reference_Manual_Complete2_Optimized.pdf).
- 598 35. Flores, M.V.; Ertola, R.J.; Voget, C.E. Effect of monovalent cations on the stability
599 and activity of *Kluyveromyces lactis* β -galactosidase. *Lebensm. -Wiss. u. -Technol.*
600 **1996**, 29, 503-506.
- 601 36. Garman, J.; Coolbear, T.; Smart, J. The effect of cations on the hydrolysis of lactose
602 and the transferase reactions catalysed by β -galactosidase from six strains of lactic acid
603 bacteria. *Appl. Microbiol. Biotechnol.* **1996**, 46, 22-27.
- 604 37. Rastall, R. Functional oligosaccharides: application and manufacture. *Annu. Rev.*
605 *Food Sci. Technol.* **2010**, 1, 305-339.
- 606 38. Gibson, G.R.; Probert, H.M.; Loo, J.V.; Rastall, R.A.; Roberfroid, M.B. Dietary
607 modulation of the human colonic microbiota: updating the concept of prebiotics. *Nutr.*
608 *Res. Rev.* **2004**, 17, 259-275.
- 609 39. Valette, P.; Pelenc, V.; Djouzi, Z.; Andrieux, C.; Paul, F.; Monsan, P.; Szylit, O.
610 Bioavailability of new synthesised glucooligosaccharides in the intestinal tract of
611 gnotobiotic rats. *J. Sci. Food Agric.* **1993**, 62, 121-127.
- 612 40. Nakada, T.; Nishimoto, T.; Chaen, H.; Fukuda, S. Kojioligosaccharides: Application
613 of kojibiose phosphorylase on the formation of various kojioligosaccharides.
614 Oligosaccharides in Food and Agriculture. In *Oligosaccharides in Food and*
615 *Agriculture. ACS Symposium Series*; Eggleston, G.; Côté, G. L., Eds.; American
616 Chemical Society: Washington, DC, 2003; Vol. 849, pp. 104-117.
- 617 41. Sanz, M.L.; Gibson, G.R.; Rastall, R.A. Influence of disaccharide structure on
618 prebiotic selectivity in vitro. *J. Agric. Food Chem.* **2005**, 53, 5192-5199.

619 **Figure captions**

620

621 **Figure 1.** HPLC-RID profiles of transglycosylation reactions based on sucrose:lactose
622 and sucrose:whey permeate mixtures catalyzed by dextransucrase from *Leuconostoc*
623 *mesenteroides* B-512F (0.8 U mL⁻¹) at 30 °C, in 20 mM sodium acetate buffer at pH
624 5.2, for 0 and 24 hours. **A)** sucrose:lactose 45%:20%; **B)** sucrose:lactose 30%:30%; **C)**
625 sucrose:whey permeate 2 (equivalent to a concentration of sucrose:lactose of
626 30%:30%).

627 Labelled peaks are as follows: 1 (fructose); 2 (glucose); 3 (sucrose); 4 (leucrose); 5
628 (lactose); 6 (minor acceptor-reaction products); 7 (2- α -D-glucopyranosyl-lactose).

629

630 **Figure 2.** Concentrations of sucrose, lactose, leucrose and 2- α -D-glucopyranosyl-
631 lactose upon transglycosylation reactions based on sucrose:lactose and sucrose:whey
632 permeate mixtures catalyzed by dextransucrase from *Leuconostoc mesenteroides* B-
633 512F (0.8 U mL⁻¹) at 30 °C in 20 mM sodium acetate buffer at pH 5.2. **A)**
634 sucrose:lactose 45%:20%; **B)** sucrose:lactose 30%:30%; **C)** sucrose:whey permeate 1
635 (equivalent to a concentration of sucrose:lactose of 30%:30%); **D)** sucrose:whey
636 permeate 2 (equivalent to a concentration of sucrose:lactose of 30%:30%). Sucrose
637 (—○—), lactose (—△—), leucrose (—□—), 2- α -D-glucopyranosyl-lactose (—◆—).
638 Vertical bars represent standard deviations ($n = 3$).

639

640 **Figure 3.** HPAEC-PAD profile of transglycosylation reaction based on sucrose:lactose
641 (30%:30%) catalyzed by dextransucrase from *Leuconostoc mesenteroides* B-512F (0.8
642 U mL⁻¹) at 30 °C, in 20 mM sodium acetate buffer at pH 5.2, for 24 hours. Labelled
643 peaks are as follows: 1 (glucose), 2 (fructose), 3 (leucrose), 4 (lactose), 5 (2- α -D-

644 glucopyranosyl-lactose). (*) The inset shows a zoom area of the eluted minor acceptor-
645 reaction products of degree of polymerization above 3.

646

647 **Figure 4.** 2D TOCSY NMR spectrum with water suppression corresponding to 2- α -D-
648 glucopyranosyl-lactose obtained at 500 MHz in D₂O. **A)** Contour plot of the sugar
649 region. **B)** Horizontal traces corresponding to the resonances of anomeric protons of
650 galacto- and gluco-residues.

Table 1. ^1H (500 MHz) and ^{13}C (125 MHz) NMR spectral data for 2- α -D-glucopyranosyl-lactose. Chemical shift (δ , ppm) and coupling constants (J in Hz, in parenthesis).

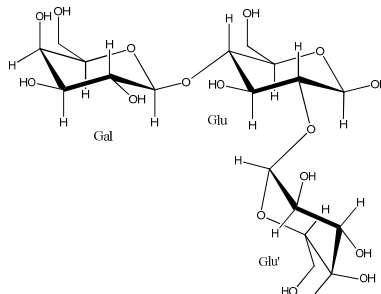
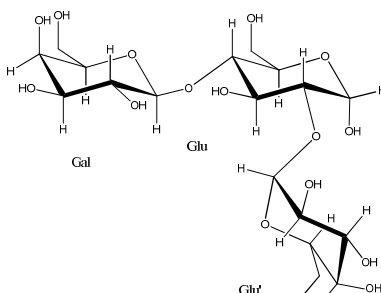
Isomer	Position	Gal		Glu		Glu'	
		δ_{H}	δ_{C}	δ_{H}	δ_{C}	δ_{H}	δ_{C}
β anomer 	1	4.33 (6.3)	103.9 9	4.70 (8.0)	97.07	5.24 (3.9)	98.81
	2	3.43	72.04	3.31	79.40	3.43	72.50
	3	3.54	73.62	3.63	74.12	3.64	73.89
	4	3.81	69.62	3.56	79.73	3.35	70.33
	5	3.60	76.37	3.48	75.67	3.95	72.58
	6	3.65	62.07	3.70- 3.84	61.35	3.68	61.24
α anomer 	1	4.35 (6.1)	103.9 0	5.32 (3.5)	90.23	4.98 (3.8)	97.46
	2	3.45	72.04	3.58	76.64	3.43	72.44
	3	3.55	73.62	3.84	70.90	3.68	73.84
	4	3.81	69.62	3.59	79.43	3.34	70.43
	5	3.60	76.37	3.86	71.03	3.86	72.78
	6	3.65	62.07	3.70- 3.84	61.04	3.86	61.24

Table 2. Physical-chemical characterization of industrial cheese whey permeates (WP1 and WP2).

	WP1	WP2
Dry matter (DM)	94.2 ± 0.0	95.5 ± 0.2
Lactose (% w/w DM)	89.9	81.6
Protein (% w/w DM)	0.1 ± 0.0	Tr*
Mineral (mg/g DM)	24.6	47.0
Main elements (mg/g DM)		
Sodium	3.2	6.8
Magnesium	1.0	1.3
Potassium	6.3	26.5
Calcium	9.8	5.5
Phosphorus	4.2	7.0
Minor elements (µg/g DM)		
Boron	17	19
Zinc	5	1
Bromine	9	35
Rubidium	7	24
Strontium	4	2
Iodine	Tr*	3
Titanium	n.d.**	6
Barium	n.d.**	9
pH	5.4	6.6

*Tr: Traces

**n.d: not detected

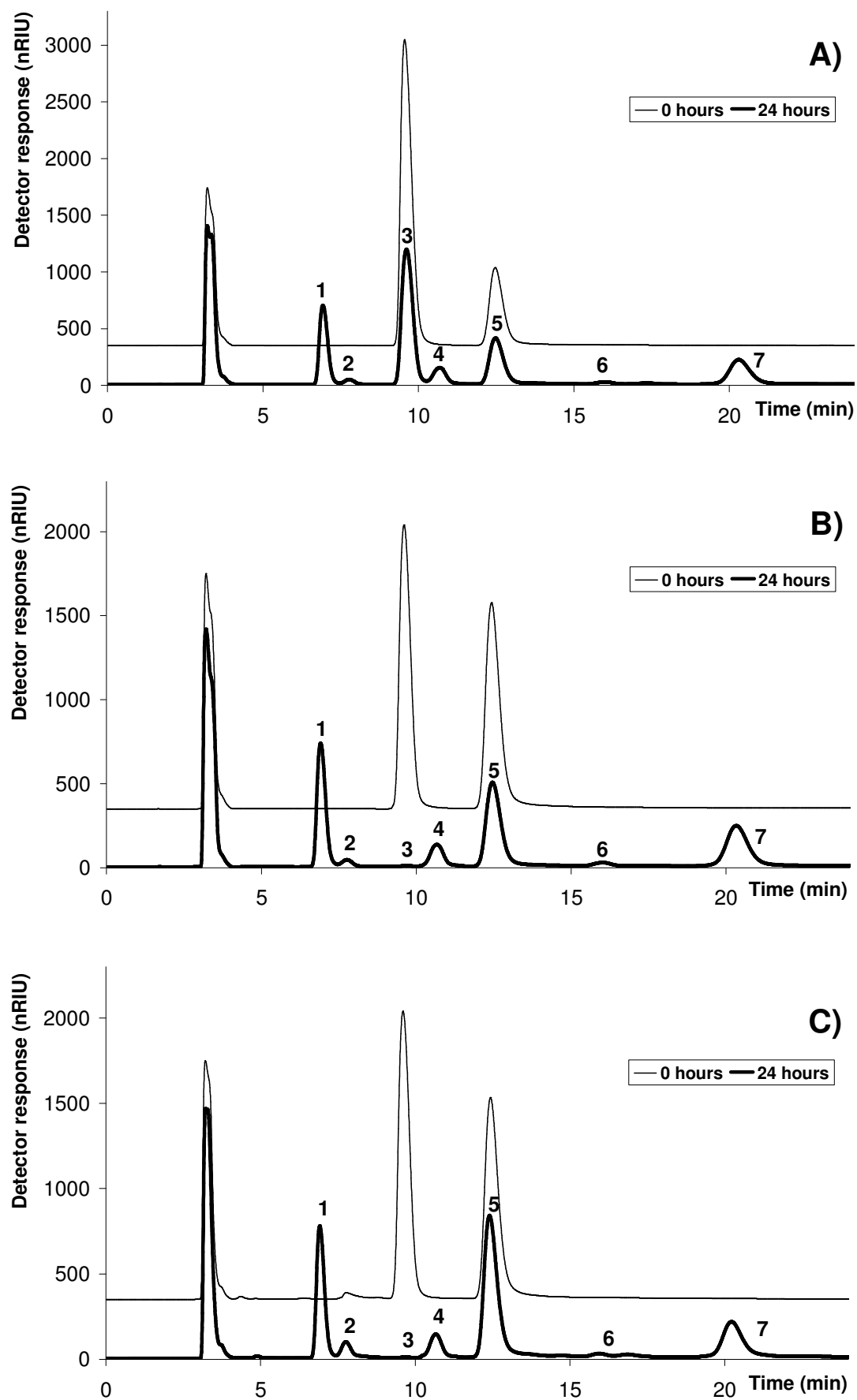
Figure 1. *Díez-Municio et al.*

Figure 2. Díez-Municio et al.

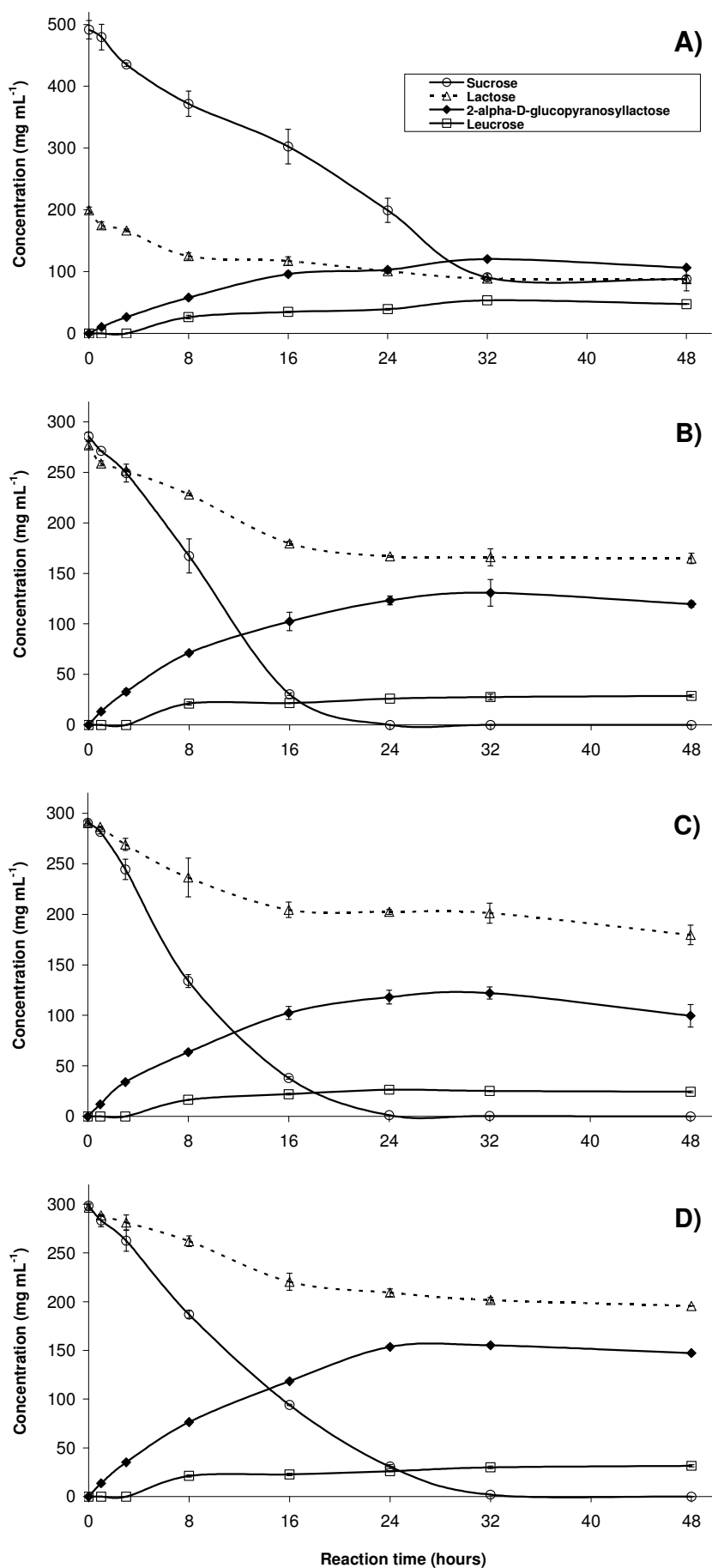


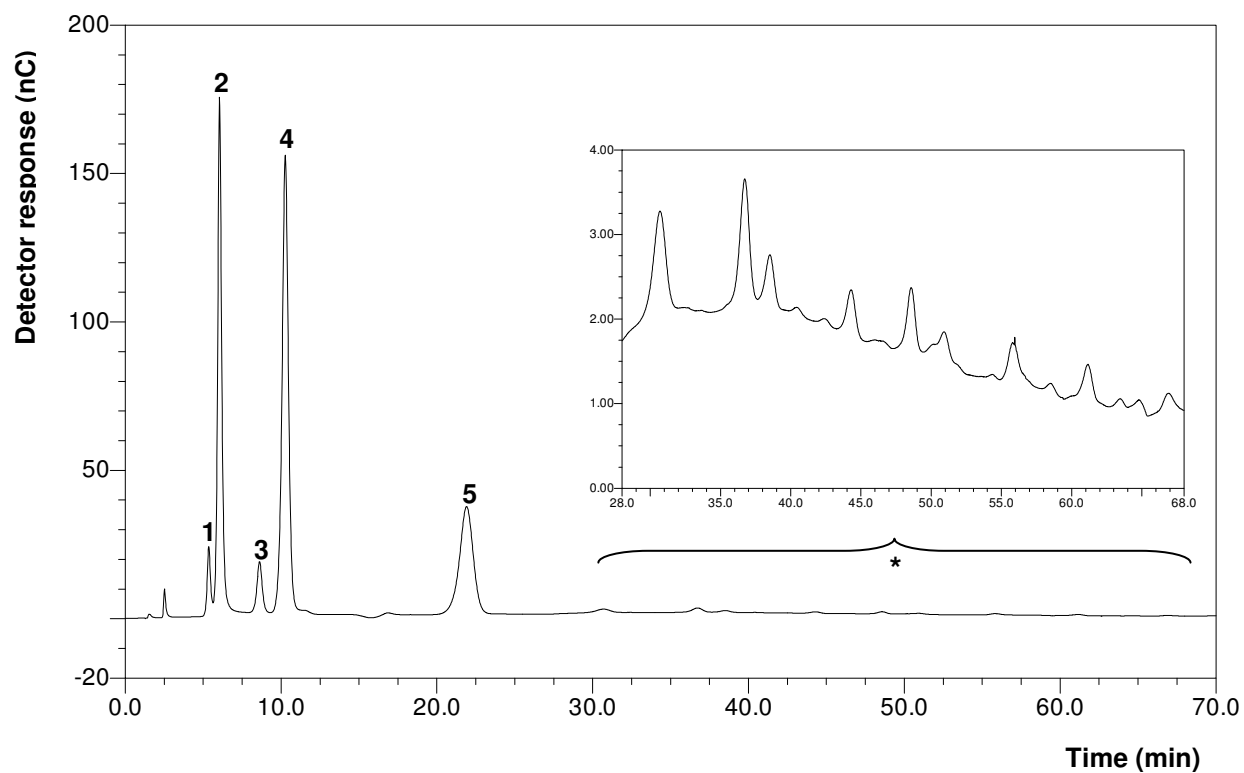
Figure 3. *Díez-Municio et al.*

Figure 4. Díez-Municio *et al.*