1	A study of influential factors on oligosaccharide formation by fructosyltransferase
2	activity during stachyose hydrolysis by Pectinex Ultra SP L
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19 ABSTRACT

20 Enzyme-catalysed transglycosylation has gained increased interest as an approach to 21 obtain novel prebiotic oligosaccharides. Influence of reaction conditions for 22 oligosaccharide synthesis from stachyose by a commercial enzymatic extract from 23 Aspergillus aculeatus (Pectinex Ultra SP-L) was studied. During this reaction, 24 oligosaccharides with degree of polymerization (DP) ranging from 3 to 8 were detected 25 by gas chromatography with flame ionization detection (GC-FID) and MALDI-TOF-26 MS. Galactosyl-melibiose (DP₃) was synthesized as a result of fructosidase activity 27 while fructosyl-stachyose (DP₅) and diffuctrosyl-stachyose (DP₆) were formed as 28 consequence of fructosyltransferase activity of Pectinex. The optimal reaction 29 conditions for the synthesis of penta- and hexasaccharides were 60 °C, pH 5.5, 600 30 mg/mL stachyose and 34 U/mL of enzyme. Composition on oligosaccharides of 31 synthesis mixtures was noticeable affected by reactions conditions. Reaction time 32 played an important role on oligosaccharide mixtures composition constituted by 33 20.05% pentasaccharides, 0.72% hexasaccharides, 54.82% stachyose, 20.75% 34 and 1.03% monosaccharides after 1 h and galactosyl-melibiose 16.27% 35 pentasaccharides, 3.83% hexasaccharides, 27.26% stachyose, 43.61% galactosyl-36 melibiose and 2.35% monosaccharides after 3 h. In conclusion, stachyose could be used 37 as a substrate for the enzymatic synthesis of new oligosaccharides that may open new 38 opportunities in the development of future prebiotics.

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42 Keywords: α-galactosides, stachyose, Pectinex Ultra SP-L, transfructosylation,
43 oligosaccharide synthesis

44 **1. Introduction**

45 The consumption of prebiotic carbohydrates has gained increased interest due to 46 their recognition as agents inducing beneficial physiological effects in the colon and in 47 extra-intestinal compartments or reduces the risk of associated intestinal and systemic 48 pathologies (Roberfroid et al., 2010). Prebiotic carbohydrates may be fermented in 49 different parts of digestive tract. Rapidly fermented prebiotics stimulate growth and 50 activity of beneficial bacteria in the proximal colon whereas slowly fermented 51 prebiotics may reach the distal colon where most of chronic diseases (e.g. colorectal 52 cancer and ulcerative colitis) take place (Sanz, Cote, Gibson, & Rastall, 2006; Sanz, 53 Gibson, & Rastall, 2005). Consequently, there is greater interest in finding prebiotics 54 that can persist to more distal regions of the colon. (Hughes et al., 2007; Martinez-55 Villaluenga et al., 2008; Cardelle-Cobas, Martínez-Villaluenga, Villamiel, Olano & 56 Corzo 2008).

57 In addition, monosaccharide moieties and linkage type may influence the 58 oligosaccharide fermentation rate by colonic microbiota (Sanz et al., 2006). Currently, 59 the development of future prebiotic targets the modulation of microbial fermentation 60 along the gastrointestinal tract (Ouwehand, Derrien, de Vos, Tiihonen, & Rautonen, 61 2005). Future prebiotics could be mixtures of oligosaccharides that have different 62 effects in different parts of the gastrointestinal tract. Besides colon, oral cavity and the 63 small intestine are also considered to be potential prebiotic targets (Ouwehand et al., 64 2005).

65 α -Galactosides (raffinose, stachyose, verbascose and ajugose) are sucrose-based 66 oligosaccharides found more abundantly in grain legumes (Martinez-Villaluenga, Frias, 67 & Vidal-Valverde, 2008a). These carbohydrates derived from sucrose containing 1-4 68 units of galactose linked by α -(1,6) linkages to the glucose moiety. These

oligosaccharides are resistant to digestion promoting the growth of beneficial bacteria in
the colon (Van den Broek, Hinz, Beldman, Doeswijk-Voragen, Vincken, & Voragen,
2005).

72 In vitro assays have shown the ability of Bifidobacterium and Lactobacillus 73 strains to grow using α -galactosides as carbon source, due to the production of α -74 galactosidase (Trojanova, Vlkova, Rada, & Marcunek, 2006; Farnworth, Mainville, 75 Desjardins, Gardner, Fliss, & Champagne, 2007; Martinez-Villaluenga, & Gomez, 76 2007). Today, α -galactosides are obtained on industrial scale as a by-product from the 77 production of soy protein isolates and soy protein concentrates (Martinez-Villaluenga, 78 et al., 2008a). The recovery of α -galactosides from soy by-products could be an 79 interesting alternative for their utilization as raw material in the manufacture of future 80 prebiotics.

Transglycosylation reactions catalysed by enzymes are being used for the 81 82 production of oligosaccharides mixtures with a specific composition. Production of 83 galactooligosaccharides mixtures from lactose or lactulose using galactosyltransferases 84 from different sources have been carried out (Chockchaisawasdee, Athanasopoulos, Niranjan, & Rastall, 2005; Cardelle-Cobas, Villamiel, Olano, & Corzo, 2008; Martínez-85 86 Villaluenga, Cardelle-Cobas, Corzo, Olano, & Villamiel, 2008b). In addition, 87 fructooligosaccharides mixtures were produced from sucrose by fructosyltransferases 88 from bacteria or moulds (Hang & Woodams, 1995; Yun, 1996; Sangeetha, Ramesh, & 89 Prapulla, 2005; Ghazi, Fernandez-Arrojo, Gomez de Segura, Alcalde, Plou, & 90 Ballesteros, 2006). Since α -galactosides contain a fructosyl end, fructosyltransferases 91 may transfer the fructosyl moiety to a suitable acceptor to give rise new 92 oligosaccharides. Purified fructosyltransferase from Aspergillus niger has been used to 93 catalyse the synthesis of new oligosaccharides from raffinose (Uhm, Baek, Jhon, Kim,

94 Hwang, & Ryu, 1999). The relatively inexpensive commercial enzyme preparation 95 Pectinex Ultra SP-L, produced by Aspergillus aculeatus, was shown to contain 96 fructosyltransferase activity (Hang & Woodams, 1995; Tanriseven & Aslan, 2005; 97 Ghazi et al., 2006; Montilla, Corzo, Olano, & Jimeno, 2009); therefore, it could be used 98 as catalyst in large-scale production of new oligosaccharides. Therefore, the objective of 99 this work has been to investigate in more detail the fructosyltransferase activity from 100 Aspergillus aculeatus of commercial enzymatic preparation Pectinex Ultra SP-L over 101 stachyose hydrolysis.

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103 **2. Material and methods**

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105 2.1. Chemicals

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107 D-galactose, D-glucose, D-fructose, melibiose, raffinose, stachyose and phenyl-108 β -glucoside were purchased from Sigma (St. Louis, MO, USA). The commercial 109 enzyme preparation from *Aspergillus aculeatus*, Pectinex Ultra SP-L, containing 110 fructosyltransferase activity was a generous gift from Novozymes (Dittingen, 111 Switzerland).

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- 113 2.2. Determination of enzymatic activity
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115 Fructosyltransferase activity of Pectinex was measured using sucrose as 116 substrate. The activity was assayed at 60 °C using sucrose at 300g/L in 0.1 M sodium 117 acetate buffer (pH 5.5). Aliquots (0.1 mL) were withdrawn at different times. The 118 reaction was stopped by adding 10 μ L of acetic acid 0.1 N. The amount of fructose and 119 glucose released was determined by gas chromatography (GC-FID) using the method 120 described below. Fructosyltransferase activity was calculated as the difference between 121 the amount of free glucose and fructose which indicated the amount of fructose 122 involved in the transfructosylation reaction. The fructosyltransferase activity was 400 123 U, where 1 unit (U) is defined as the amount of enzyme transferring 1 µmol of fructose 124 per minute per mL under assayed conditions. Enzyme activity measurements were 125 performed in triplicate and the experimental error (RSD relative standard deviation) was 126 lower than 5%.

127 The soluble protein concentration in the enzymatic preparation was determined 128 using the bicinchoninic acid (BCA) assay (Smith et al., 1985). Bovine serum albumin 129 (BSA) was used as standard. Specific activity was defined as U/mg protein. The soluble 130 protein concentration in the enzyme preparation was 82 mg/mL. Therefore, the enzyme 131 expressed a fructosyltransferase specific activity of 4.9 U/mg soluble protein.

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133 2.3. Enzymatic synthesis of oligosaccharides

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135 Enzymatic synthesis of oligosaccharides from stachyose using Pectinex was 136 carried out under different reaction conditions such as temperature (50, 60 and 70 °C), 137 pH (3.5; 4.5; 5.5; 6.5 and 7.5), stachyose concentration (100, 300 and 600 g/L), 138 enzyme concentration (17, 34 and 78 U/mL), and time (0.5, 1, 3, 6 and 24 h). Reactions 139 were performed at a final volume of 1.5 mL in microtubes incubated in an orbital shaker 140 at 300 rpm. Aliquots (120 µL) were withdrawn from the reaction mixture at the 141 different times and immediately immersed in boiling water for 5 min to inactivate the enzyme. Samples were stored at -18° C for subsequent analysis. Experiments were 142 143 carried out in duplicate.

144 The amount of remaining stachyose and the yield of oligosaccharides in the 145 reaction mixtures were expressed as weight percentage of total carbohydrate content.

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147 2.4. Purification and characterization of reaction mixtures

148 Purification was performed following the method described by Morales, Sanz, 149 Olano, & Corzo, (2006). Briefly, a total of 2.5 mL of reaction mixture, containing 1.5 g 150 of carbohydrates was dissolved in 300 mL of water and stirred for 30 min with 9 g of 151 activated charcoal Darco G60, 100 mesh (Sigma, St. Louis, MO) to remove mono and 152 disaccharides. This mixture was vacuum-filtered through Whatman No. 1 filter paper, 153 and activated charcoal was washed with 50 mL of water. The oligosaccharides adsorbed 154 onto the activated charcoal were extracted by stirring for 30 min with 300 mL 155 ethanol/water solution (1:1, v:v) and then vacuum-filtered. The ethanol /water solution 156 was evaporated under vacuum at 30 °C. Sample was dissolved in 5 mL of deionized 157 water and filtered through 0.22 µm filters (Millipore Corp., Bedford, MA) for further 158 characterization by mass-spectrometry.

159 The enriched fraction containing the main synthesis products was characterized 160 by Matrix-Assisted Laser Desorption/Ionisation Time-of-Flight Mass Spectrometry 161 (MALDI-TOF-MS) on a Voyager DE-PRO mass spectrometer (Applied Biosystems, 162 Foster City, CA) equipped with a pulsed nitrogen laser ($\lambda = 337$ nm, 3 ns pulse width, 163 and 3 Hz frequency) and a delayed extraction ion source. Ions generated by laser 164 desorption were introduced into a time of flight analyser (1.3 m flight path) with an 165 acceleration voltage of 20 kV, 74% grid voltage, 0.001% ion guide wire voltage, and a 166 delay time of 300 ns in the reflector positive ion mode. Mass spectra were obtained over 167 the m/z range 100-1500. External mass calibration was applied using the monoisotopic $[M + H]^+$ values of des-Arg¹ bradykinin and angiotensin I of the Calibration Mixture 1, 168

169 Sequazyme Peptide Mass Standards Kit; Applied Biosystems. 2,5-dihydroxybenzoic 170 acid (>98%, Fluka) at 10 mg/mL in water was used as matrix. Sample was diluted 100 171 times in water, and mixed with the matrix at a ratio of 1:4 (v:v). One μ L of this solution 172 was spotted onto a flat stainless-steel sample plate and dried in air before analysis.

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174 2.5. Carbohydrate analysis by gas chromatography (GC-FID)

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176 Trimethyl silvlated oximes (TMS-oxime) of mono- di and oligosaccharides were 177 analysed by GC following the method of Montilla, van de Lagemaat, Olano, & del 178 Castillo (2006). GC analysis was performed with a gas chromatograph (3800GC, 179 Varian, California, USA) equipped with a flame ionization detector (FID). The 180 trimethylsilyl oximes were separated using a 8 m x 0.25 mm x 0.25 µm film fused silica 181 capillary column coated with CP-SIL 5CB (methyl silicone from Chrompack, 182 Middelburg, The Netherlands), giving rise to two peaks, corresponding to the syn (E) 183 and anti (Z) isomers. The carrier gas (nitrogen) flow rate was 1.1 mL/min. Injector 184 temperature was 280 °C, detector temperature was 340 °C and the oven temperature was 185 programmed from 150 °C to 165 °C at 3 °C/min, then at 5 °C/min to 340 °C and keeping 186 this temperature for 10 min. Injections were made in the split mode (1:10). Data 187 acquisition and integration was done using HP ChemStations software (Hewlett-188 Packard, Wilmington, USA).

The oximes were formed following the method of Brobst & Lott (1966). A
volume of 7 to 40 μl of sample constituted of approximately 4 mg of sugars, was added
to 0.4 mL of internal standard (IS) solution containing 0.5 mg/ml phenyl-β-glucoside.
The mixture was dried at 38-40 °C in a rotary evaporator (Büchi Labortechnik AG,
Flawil, Switzerland). Sugar oximes were formed by adding 200 μL hydroxylamine

194 chloride (2.5%) in pyridine and heating the mixture at 70 °C for 30 min. Subsequently, 195 the oximes obtained in this step were silylated with hexamethyldisilazane (200 μ L) and 196 trifluoroacetic acid (20 μ L) and kept at 50 °C for 30 min. Reaction mixtures were 197 centrifuged at 7000*g* for 5 min at 5 °C. Supernatants were injected in the GC or stored 198 at 4 °C prior to analysis.

Quantitative analysis was carried out by the internal standard method. Response
factors were calculated using a triplicate analysis of 6 standard solutions (galactose,
glucose, fructose, sucrose, melibiose, raffinose and stachyose) at a concentration range
from 0.06 to 2 mg/ml and 0.06 to 4 mg/ml for stachyose.

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204 **3. Results and discussion**

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206 The fructosidase, galactosidase and fructosyltransferase activities of Pectinex 207 resulted in the production of melibiose (α -D-Galp-($1\rightarrow 6$)- α -D-Glcp), raffinose (α -D-208 Galp- $(1\rightarrow 6)$ - α -D-Glcp- $(1\rightarrow 2)$ - β -D-Fruf), 6' galactosyl melibiose (α -D-Galp- $(1\rightarrow 6)$ - α -D-Galp- $(1 \rightarrow 6)$ - α -D-Glcp), fructosyl stachvose (DP5) (1^F fructofuranosyl stachvose, α -209 210 D-Galp- $(1\rightarrow 6)$ - α -D-Galp- $(1\rightarrow 6)$ - α -D-Glcp- $(1\rightarrow 2)$ - β -D-Fruf- $(1\rightarrow 2)$ - β -D-Fruf) and diffuctosyl stachyose (DP6) $(1^{F} (fructofuranosyl)_{2} \text{ stachyose } (\alpha-D-Galp-(1\rightarrow 6)-\alpha-D-$ 211 212 Galp- $(1\rightarrow 6)$ - α -D-Glcp- $(1\rightarrow 2)$ - β -D-Fruf- $(1\rightarrow 2)$ - β -D-Fruf- $(1\rightarrow 2)$ - β -D-Fruf) as it has 213 been previously reported (Montilla et al., 2009). Besides, MALTI-TOF-MS analysis of 214 oligosaccharide-enriched fractions allowed the detection of oligosaccharides with DP7 215 (m/z, 1175.4 and 1191.3) and DP8 (m/z, 1338.5) (Fig. 1) whereas previous studies have 216 only reported the formation of oligosaccharides DP3 and DP5 from stachyose by 217 fructosyltransferase from Aspergillus niger (Uhm et al., 1999)

219 Several reports have demonstrated that glycosidic linkages and degree of 220 polymerization of oligosaccharides contribute toward the selectivity of fermentation by 221 beneficial bacteria (Rowland, & Tanaka, 1993; Sanz et al., 2005; 2006). In addition, a 222 wide molecular weight range could give rise to modulation of microbial fermentation. 223 Higher molecular oligosaccharides may be slowly fermented exhibiting higher colonic 224 persistence than low molecular weight carbohydrates, reaching the most distal regions 225 where most of intestinal disorders are encountered (Sanz et al., 2006). On the basis of 226 this information, the production of α -galactosides mixtures from stachyose composed of 227 carbohydrates with DP from 3 to 8 may represent an opportunity in the development of 228 future prebiotics. 229 230 3.1. Study of the formation of oligosaccharides from hydrolysis of stachyose 231 232 A series of experiments were performed in which temperature, pH and sustrate 233 and enzyme concentrations were varied to select the optimum conditions for the tri- and 234 oligosaccharides syntesis. 235 236 3.1.1. Effect of temperature 237 To determine the influence of temperature on oligosaccharide synthesis, reaction 238 was performed in a temperature range of 50-70 °C, 300 mg/mL of stachyose, 34 U/mL 239 of enzyme and pH 5.5, for 24 h (Fig. 2). The loss of stachyose (Fig. 2a) ncreased with 240 temperature from 50° to 60°C, but declined with further increase of temperature to 70°C 241 This fact could be attributed to a thermal denaturation of enzyme at 70 °C. 242 During the first six hours of the reaction fructose release (Fig. 2b) was favoured

243 at 60 °C, however, after twenty four hours, when most of the stachyose was consumed,

the amount of fructose reached at 50°C was similar to that observed at 60°C Galactosylmelibiose production (Fig. 2c) reached maximum yields after 24 h either at 50 °C or 60
°C (58.9% and 62.9%, respectively).

247 Transferase/hydrolase ratio of the enzyme was optimal at 60 °C, being observed 248 the highest DP5 oligosaccharide yield (15.3%) after 3 h (Fig. 2d). It is noteworthy a 249 sharply decrease of this pentasaccharide at 50 and 60 °C after reaching maximum yield 250 while larger amounts of this α-galactoside remained after 24 h at 70 °C. DP6 251 oligosaccharide production was only possible at 60 °C, reaching the highest 252 concentration (1.8%) after 6 h. Our results are in agreement with previous works which 253 reported that the optimal temperature for fructooligosaccharides production from 254 sucrose by fructosyltransferase activity of Pectinex was around 60 °C (Tanriseven & 255 Aslan, 2005; Ghazi, Fernandez-Arrojo, Garcia-Arellano, Ferrer, Ballesteros, & Plou, 256 2007).

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258 *3.1.2. Effect of pH*

Different experiments were carried out at pH values from 3.5 to 7.5 at 60 °C using reactions mixtures of stachyose (300 mg/mL) and 34 U/mL of enzyme. In assays carried out at pH 3.5 and 7.5 hardly any stachyose hydrolysis was observed (data not shown). The highest rate of stachyose decrease (Fig. 3a), fructose release (Fig. 3b) and galactosyl-melibiose formation (Fig. 3c) were attained at pH 5.5. At pH 4.5 the lowest fructosidase activity was observed.

The maximum formation of DP5 (Fig. 3d) was observed at pHs 5.5 and 6.5 after three hours ($\sim 16\%$) and then, at pH 6.5 slowly decreased down to ($\sim 12\%$) after twenty four hours. However, a markedly decrease to ($\sim 2\%$) was observed at pH 5.5 probably due to a higher fructosidase activity of the enzyme under these conditions.

The highest amount of DP6- formed was 1.8%, at pH 5.5 after 6 h of reaction whereas yields at pH 4.5 and 6.5 were negligible. These results are consistent with previous studies of Ghazi et al. (2007) showing that the optimal activity/stability of purified fructosyltransferase from Pectinex was in the pH range 5.0-6.0; similar 1y to other fructosyltransferases derived from fungi (Hang, & Woodams, 1996; Rezende, & Felix, 1997).

275 *3.1.3. Effect of enzyme concentration*

276 The effect of enzyme concentration (17, 34 and 68 U/mL) was studied using 277 reaction mixtures containing 300 mg/mL stachyose at pH 5.5 and at 60 °C (Fig. 4). Loss 278 of stachyose and formation of galactosyl-melibiose noticeably increased when enzyme 279 concentration was raised up to 34 U/mL while a further increase of the enzyme 280 concentration did not result in a significant improve of stachyose conversion and 281 trisaccharide yields (Fig. 4a and 4c). Regarding DP5 formation, enzyme concentration 282 did not exert any influence on maximum yields attained, although formation rate of this 283 pentasaccharide increased with enzyme concentration (Fig. 4d). DP6 yields decreased 284 with enzyme concentration, the highest yield (2.1%) was observed at enzyme 285 concentration of 17 U/mL after 24 h (Fig. 4d). At the same with stachyose, the most 286 important effect observed was an increase of hydrolysis rate with enzyme concentration. 287 Also, hydrolysis rate of the DP5 and DP6 formed during the first three to six hours 288 increased with the enzyme concentration (Fig. 4d). This was consistent with the 289 monosaccharide concentrations observed (Fig. 4b). The highest enzyme concentration 290 tested (68 U/mL) gave rise to the highest fructose (13.3%) and galactose+glucose 291 (4.8%) concentrations after 24 h of reaction. However, in all conditions it was observed 292 that galactosyl-melibiose was stable to hydrolysis; the highest melibiose formation was 293 only 3.4% after 24h with 68 U/mL of enzyme concentration.

295 *3.1.4. Effect of initial stachyose concentration*

296 To investigate the influence of substrate concentration on oligosaccharide 297 synthesis, reaction was performed using a range of stachyose concentration from 100 to 298 600 mg/mL (Fig. 5). Reactions were performed at 60 °C, pH 5.5 and 34 U/mL of 299 enzyme for 24 h. Fig. 5a shows that at the higher initial substrate concentration the 300 higher rate of stachyose conversion was reached. The fructosidase activity of Pectinex 301 was favoured with low stachyose concentrations (100 mg/mL) and longer reaction time, 302 thus increasing amounts of free fructose was released reaching up to 13.8% after 24 h of 303 reaction (Fig. 5b).

304 Formation of galactosyl-melibiose increased with higher initial stachyose 305 concentration up to 6 h (Fig. 5c), and the highest galactosyl-melibiose yield (67.5%) 306 was found with 100 mg/mL of initial stachyose concentration after 24 h of reaction. The 307 galactosyl-melibiose yield obtained by Pectinex using stachyose as substrate was much 308 higher than that obtained by Van Laere et al., (1999) who achieved a 33 % using α -309 galactosidase from Bifidobacterium adolescentis and 300 mM melibiose. Tzortzis, Jay, 310 Baillon, Gibson, & Rastall, (2003) and Goulas, Goulas, Tzortzis, & Gibson, 2009 311 obtained a 26% and 21% of galactosyl-melibiose using α -galactosidase from 312 Lactobacillus reuteri and 670 mM melibiose and α -galactosidase from Bifidobacterium 313 bifidum and 1170 mM melibiose, respectively. Galactosyl-melibiose is an interesting 314 compound with prebiotic and symbiotic properties with Lactobacillus acidophilus and 315 L. reuteri. Prebiotics properties has been found in synthesised galactosyl melibiose 316 mixtures which produced a higher increase in bifidobacteria and lactobacilli population 317 and decrease of clostridia and *E. coli* population than produced by FOS, melibiose and 318 raffinose used as reference carbohydrates (Tzortzis, Goulas, Baillon, Gibson, & Rastall,
319 2004).

320 Transfructosylation reaction-mainly occurred at reaction times less than three 321 hours and increased with initial substrate concentrations up to 600 mg/mL (Fig. 5d). 322 The highest yields of transfructosylation products DP5 (20.1%) and DP6 (3.8%), were 323 found at the highest initial stachyose concentration (600 mg/mL) after 1 h and 3 h, 324 respectively. Our research group used previously this concentration to obtain and 325 charaterize DP5 and DP6 compounds (Montilla et al., 2009). These results are 326 consistent with previous studies showing that high substrate concentration exerts a 327 noticeable influence on the formation of stachyose-derived oligosaccharides by 328 fructosyltransferases due to an increase in transferase/hydrolysis ratio (Mahoney, 1998; 329 Cardelle-Cobas, et al. 2008). Ghazi et al. (2007) reported that the fructosyltransferase 330 activity in a purified enzyme from A. aculeatus is approximately 20-fold higher than 331 hydrolysis activity at sucrose concentrations greater than 342 g/L.

332

333 4. Conclusions

334 These results shown the noticeable effect of reaction conditions on the 335 oligosaccharide formation during the enzymatic treatment of stachyose with Pectinex 336 Ultra SP-L. The optimal reaction conditions for the synthesis of penta- and 337 hexasaccharides were 60 °C, pH 5.5, 600 mg/mL stachyose and 34 U/mL of enzyme. 338 Under these conditions the time of reaction may noticeably influence the composition 339 of α -galactoside mixtures, thus, 0.72% hexasaccharides, 20.05% pentasaccharides, 340 54.82% stachyose, 20.75% galactosyl-melibiose and 1.03% monosaccharides was found 341 after 1 h of reaction; and 3.83% hexasaccharides, 16.27% pentasaccharides, 27.26% 342 stachyose, 43.61% galactosyl-melibiose and 2.35% monosaccharides after 3 h of time

course reaction. However, to obtain the maximum yield of galactosyl-melibiose
(67.5%), the assays should be carried out at 60 °C and pH 5.5, using 100 mg/mL
stachyose and 34 U/mL of enzyme during 24h.

 α -Galactosides which are industrially available in large amounts as a by-product from the production of soy protein isolate seems to be a promising raw material for the production of new oligosaccharides through hydrolysis and transglycosylation using a GRAS commercial enzyme preparation. Further research is warranted to evaluate the prebiotic properties of these new α -galactosides mixtures that may be used to influence the microbial composition in the distal colon, where most of gut disorders occur.

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353 Acknowledgements

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We are grateful to Ramiro Martinez (Novozymes A/S, Spain) for providing us with Pectinex Ultra SP-L and Plácido Galindo for his help in acquiring the MALDI-TOF-MS spectral data. This work has been financed under a R+D program of the Spanish Ministry of Science and Innovation Science, Projects AGL-2008-00941/ALI and Consolider Ingenio 2010 (FUN-C-FOOD) CSD 2007-00063; R + D program of the Comunidad de Madrid, Project ALIBIRD P2009/AGR-1469 and as R + D program of the Comunidad de Castilla-La Mancha POII10-0178-4685.

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471

472 FIGURE CAPTIONS

473 Figure 1. MALDI-TOF-MS profile of the purified fraction of oligosaccharides formed
474 during enzymatic hydrolysis of stachyose with Pectinex Ultra SP-L (DP, degree of
475 polymerisation).

476 **Figure 2.** Effect of temperature on stachyose hydrolysis and oligosaccharide production

477 during the enzymatic treatment of stachyose (300 mg/mL) with Pectinex Ultra SP-L

478 (34U/m/L) in 0.1 M sodium acetate buffer pH 5.5 at (\blacklozenge) 50 °C; (\blacksquare) 60 °C; (\blacktriangle) 70°C.

479 Figure 3. Effect of pH on stachyose hydrolysis and oligosaccharide production during

480 the enzymatic treatment of stachyose (300 mg/mL) with Pectinex Ultra SP-L (34

481 units/mL) at 60°C in 0.1 M sodium acetate buffer (\blacklozenge) pH 4.5; (\blacksquare) pH 5.5; (\blacktriangle) pH 6.5.

482 Figure 4. Effect of enzyme concentration on stachyose hydrolysis and oligosaccharide

483 production during the enzymatic treatment of stachyose (300 mg/mL) at 60°C and pH

- 484 5.5 with Pectinex Ultra SP-L (♦) 17 U/mL; (■) 34 U/mL; (▲) 68 U/mL.
- 485 Figure 5. Effect of stachyose concentration on hydrolysis and oligosaccharide
- 486 production from reaction mixtures of stachyose (\blacklozenge) 100 mg/mL; (\blacksquare) 300 mg/mL; (\blacktriangle)
- 487 600 mg/mL and Pectinex Ultra SP-L (34 U/mL), at pH 5.5 and 60 °C.
- 488 (*): Data from Montilla et al., (2009).



Figure 1. Montilla et al., 2011



Figure 2.- Montilla et al., 2011



Figure 3.- Montilla et al., 2011



Figure 4.- Montilla et al., 2011



Figure 5.- Montilla et al.,