

# In vitro fermentability of xylooligosaccharide and xylo-polysaccharide fractions with different molecular weights by human faecal bacteria

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1	In vitro fermentability of xylo-oligosaccharide and xylo-polysaccharide fractions with
2	different molecular weights by human faecal bacteria
3	
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#### 22 Abstract

23 Xylo-oligosaccharides and xylo-polysaccharides (XOS, XPS) produced by autohydrolysis 24 of the fibre from oil palm empty fruit bunches (OPEFB) were purified using gel filtration 25 chromatography to separate the XOS and XPS from the crude autohydrolysis liquor. Six mixed fractions of refined XOS and XPS with average degree of polymerisation (avDP) of 26 27 4-64 were obtained. These were characterised in terms of their composition and size by 28 HPLC, MALDI-ToF-MS (selected fractions) and carbohydrate gel electrophoresis (PACE). 29 They were assessed in batch culture fermentations using faecal inocula to determine their 30 ability to modulate the human faecal microbiota *in vitro* by measuring the bacterial growth, 31 organic acid production and the XOS assimilation profile. The gut microbiota was able to 32 utilise all the substrates and there was a link between the XOS/XPS degree of 33 polymerisation with the fermentation properties. In general, XOS/XPS preparations of 34 lower avDP promote better *Bifidobacterium* growth and organic acid production. 35 36 **Keywords** 37 Autohydrolysis; in vitro fermentation; Oil palm empty fruit bunches; Xylo-oligo and xylo-38 polysaccharides 39 40 **1. Introduction** 41 The benefits of non-digestible oligosaccharides (NDOs) in modulating the intestinal 42 and colonic microbiota that have an effect on human gut health have been well established 43 and the study of NDOs derived from plant cell walls as emerging prebiotics has raised 44 much interest. This is because plant cell walls, especially derived from cereal grains, are

45	part of our dietary fibre intake; the particular components of plant cell walls of interest in
46	the context of prebiotics are the hemicelluloses. Hemicelluloses are the second most
47	abundant class of polysaccharides available in the plant kingdom with xylan being the most
48	common. Xylans have a backbone of $\beta$ -(1 $\rightarrow$ 4) linked xylose units that are often substituted
49	with arabinose, methylated or non-methylated glucuronic acid, acetic acid or ferulic acid
50	(Ebringerová, Hromadkova & Heinze, 2005). Thus, depending on the origin of the plant
51	cell wall and treatment process, various xylo-oligosaccharides (XOS, avDP $\leq 20$ ) or xylo-
52	polysaccharides (XPS, $avDP > 20$ ) with or without branching can be obtained.
53	In this regard, plant lignocellulosic biomass generated at the agricultural field and
54	processing plant, which was once considered as waste for disposal, offers an enormous
55	potential resource as a basic feedstock for XOS production (Moure, Gullón, Domínguez &
56	Parajó, 2006). In the context of biorefining, hydrothermal treatments such as autohydrolysis
57	have been investigated as an initial step of a possible multi-stage process for the utilisation
58	of lignocellulosic materials, as it can produce soluble oligosaccharides, leaving cellulose
59	and lignin in the solid phase for other usage (Parajó, Garrote, Cruz & Domínguez, 2004).
60	The XOS obtained from autohydrolysis treatment also retain some substituents that are
61	present in the native xylan such as acetyl groups, which could have an effect on their
62	fermentability by the human intestinal microbiota (Kabel, Schols & Voragen, 2002b).
63	Earlier studies on the ability of XOS to modulate the intestinal microbiota
64	investigated linear XOS of small molecular weight with a DP around 2-3. The low
65	molecular weight XOS significantly promoted the growth of bifidobacteria and led to an
66	increase in short chain fatty acid (SCFA) production in the bacterial cultures (Crittenden et

67	al., 2002; Okazaki, Fujikawa & Matsumoto, 1990; Palframan, Gibson & Rastall, 2003a)
68	and in <i>in vivo</i> studies in humans and animals (Campbell, Fahey & Wolf, 1997; Childs et al.,
69	2014; Chung, Hsu, Ko & Chan, 2007). Pure culture studies using XOS from corn cob and
70	rice husk autohydrolysis with $DP \le 4$ also enhanced the growth of bifidobacteria despite
71	having some acetyl groups and/or uronic acid substituents (Gullón et al., 2008; Moura et
72	al., 2007). In Kabel, Kortenoeven, Schols & Voragen (2002a), a XOS preparation with
73	wider range mixed DP (DP 2-11) was used and when fermented in vitro with human faecal
74	inocula, the substrate was almost completely degraded in 20-40 h of fermentation. The
75	fermentation rate and the SFCA profiles however varied depending on the substituents that
76	were present, whereby the linear XOS and arabinose substituted XOS (AXOS) were
77	fermented faster than acetylated XOS and methylglucuronylated XOS was the slowest. The
78	bacteriology profile however was not the focus in that study, so the way the substituents
79	modulate the gut microbiota is unknown. Increases in potentially health-positive bacterial
80	groups such as Bifidobacterium spp. and Lactobacillus/Enterococcus spp. were seen with
81	high average molecular weight arabinoxylans of 66,278 and 354 kDa (Hughes, Shewry, Li,
82	Gibson, Sanz & Rastall, 2007). However, the arabinoxylans tested also significantly
83	promoted clostridial growth. Van Craeyveld (2008) in a more systematic study on the
84	influence of the average degree of polymerisation (avDP) and average degree of arabinose
85	substitution (avDAS) of XOS preparation in the cecum of rats, showed that low molecular
86	weight AXOS (avDP-avDAS of 5-0.27 and 3-0.26) increased Bifidobacterium spp.
87	significantly more than high molecular weight AXOS (avDP –avDAS of 61-0.58). On the
88	other hand, the measured branched SCFA was the lowest with avDP 61, so this could
89	potentially suppress the metabolites from protein fermentation.

90	In a previous study, results have demonstrated that it is possible to produce purified
91	XOS fractions of a variety of avDP from oil palm biomass autohydrolysis liquor (Ho et al.,
92	2014). The aim of this work was to study the effect of XOS and XPS obtained from
93	purification of autohydrolysed OPEFB at different avDP 4-64 upon the gut microbiota
94	population. The rationale for inclusion of higher avDP XOS/XPS preparations in this study
95	is they may have better persistence into the distal colon, with potential benefits to chronic
96	gut diseases.
97	
98	2. Materials and methods
99	2.1 Preparation of XOS/XPS fractions
100	The XOS/XPS preparation was according to Ho et al. (2014). Briefly, the fibre of
	The XOS/XPS preparation was according to Ho et al. (2014). Briefly, the fibre of dried oil palm empty fruit bunches (OPEFB) was subjected to non-isothermal
101	
101 102	dried oil palm empty fruit bunches (OPEFB) was subjected to non-isothermal
101 102 103	dried oil palm empty fruit bunches (OPEFB) was subjected to non-isothermal autohydrolysis treatment in a two litre capacity stainless steel reactor (Parr Instruments Co.,
101 102 103 104	dried oil palm empty fruit bunches (OPEFB) was subjected to non-isothermal autohydrolysis treatment in a two litre capacity stainless steel reactor (Parr Instruments Co., llinois, United States) with an operational temperature 210 °C and a liquid to solid ratio of 8
101 102 103 104 105	dried oil palm empty fruit bunches (OPEFB) was subjected to non-isothermal autohydrolysis treatment in a two litre capacity stainless steel reactor (Parr Instruments Co., llinois, United States) with an operational temperature 210 °C and a liquid to solid ratio of 8 (w/w). The liquor obtained from autohydrolysis treatment was filtered and purified using
101 102 103 104 105 106	dried oil palm empty fruit bunches (OPEFB) was subjected to non-isothermal autohydrolysis treatment in a two litre capacity stainless steel reactor (Parr Instruments Co., llinois, United States) with an operational temperature 210 °C and a liquid to solid ratio of 8 (w/w). The liquor obtained from autohydrolysis treatment was filtered and purified using preparative gel filtration chromatography (GFC) (Ho et al., 2014) with a BPG 100/950
100 101 102 103 104 105 106 107 108	dried oil palm empty fruit bunches (OPEFB) was subjected to non-isothermal autohydrolysis treatment in a two litre capacity stainless steel reactor (Parr Instruments Co., llinois, United States) with an operational temperature 210 °C and a liquid to solid ratio of 8 (w/w). The liquor obtained from autohydrolysis treatment was filtered and purified using preparative gel filtration chromatography (GFC) (Ho et al., 2014) with a BPG 100/950 column filled with Superdex 30 <sup>TM</sup> (Amersham Pharmacia Biotech, Uppsala, Sweden). The
101 102 103 104 105 106 107	dried oil palm empty fruit bunches (OPEFB) was subjected to non-isothermal autohydrolysis treatment in a two litre capacity stainless steel reactor (Parr Instruments Co., llinois, United States) with an operational temperature 210 °C and a liquid to solid ratio of 8 (w/w). The liquor obtained from autohydrolysis treatment was filtered and purified using preparative gel filtration chromatography (GFC) (Ho et al., 2014) with a BPG 100/950 column filled with Superdex 30 <sup>TM</sup> (Amersham Pharmacia Biotech, Uppsala, Sweden). The liquid fractions were freeze dried and then pooled together to obtained six mixed freeze-

110 2.2 Characterization of XOS/XPS fractions

111	Prior to the determination of average molar mass and chemical compositions, the
112	XOS/XPS samples were dissolved in deionised water to obtain a concentration of 10 g/L.
113	The apparent molar mass of samples was determined by high performance liquid
114	chromotography (HPLC) (Agilent 1100 series, Winnersh, UK). A size exclusion column
115	BIOSEP-SEC S2000 (Phenomenex, Cheshire, UK) was used at 30 °C with 50 mM NaNO <sub>3</sub>
116	as mobile phase at 0.7 mL/min. The eluate was detected using a refractive index (RI)
117	detector. External standards with different molecular weights, i.e. xylose,
118	maltooligosaccharides (DP 2-5) and dextrans (1-71 kDa, Sigma, Dorset, UK) were used for
119	calibration.
120	The composition of the XOS/XPS samples was assayed by HPLC to quantify free
121	monosaccharides (glucose, xylose and arabinose), aliphatic acids (acetic acid, formic acid
122	and levulinic acid) and furan derivatives (furfural and 5-hydroxymethylfurfural, HMF)
123	compounds. An Aminex HPX-87H column (BioRad, Hemel Hempstead, UK) was used at
124	50 °C with 5 mM $H_2SO_4$ as mobile phase. The monosaccharides and aliphatic acids were
125	detected with a RI detector while furfural and HMF were detected using a diode array
126	detector (DAD) at 280 nm.
127	The oligosaccharide content was determined by an indirect method using
128	quantitative acid hydrolysis; this was done by mixing the XOS/XPS sample with $H_2SO_4$
129	(72 % w/w) to obtain a final acid concentration of 4 % (w/w) and the sample was heated at
130	121 °C for 60 min to induce hydrolysis. The post hydrolysed liquor was analysed with
131	HPLC and the oligosaccharide concentration was expressed as the increase in sugar
132	monomers (Sluiter et al., 2006).

133	The total phenolic content was assayed spectrophometrically by the Folin Ciocalteu
134	method using gallic acid as standard (Singleton & Rossi, 1965).
135	
136	2.3 MALDI-Tof-MS of XOS fractions
137	MALDI-ToF-MS was used to analyse the extracted OPEFB XOS fractions in native
137	WALDI-TOF-WIS was used to analyse the extracted Of EFB XOS fractions in native
138	and permethylated form for XOS avDP 4, 7 and 14. Initial analysis of the native fractions
139	did not produce strong signals (Figure S1) so analyses were also performed with
140	permethylated fractions. Permethylation of XOS fractions was performed using the
141	NaOH/DMSO slurry method using 0.5 mL of methyl iodide (Ciucanu & Kerek, 1984).
142	Permethylated glycans were dried under a stream of nitrogen and re-dissolved in 100 $\mu$ L of
143	methanol. Five $\mu$ L of native or permethylated XOS fractions (10 mg/mL) were mixed with
144	5 $\mu L$ of 2, 5-dihydroxybenzoic acid (DHB, 10 mg/mL dissolved in 50 % MeOH with 1%
145	TFA, v/v) matrix. One $\mu$ L of native or permethylated glycans mixed with DHB matrix was
146	spotted onto a MALDI target plate and allowed to air-dry. MALDI-ToF-MS was carried
147	out using a Micromass MALDI-LR mass spectrometer (Waters, Manchester, UK) using a
148	mass acquisition between $m/z$ 450 and 3 000. The MALDI set-up was as described by
149	Marsh et al. (2011). Glycan adduct ions $[M + Na]^+$ were assigned. Experimentally
150	determined masses were interpreted using GlycoMod (ExPaSy).
151	
152	2.4 Analysis of XOS/XPS fractions by carbohydrate gel electrophoresis
153	Polysaccharide Analysis by Carbohydrate Gel Electrophoresis (PACE) was used to
154	analyse the extracted OPEFB XOS/XPS fractions. Fractions were run with and without

155	digestion with xylanase 11. Briefly, for xylanase digested samples, 200 $\mu$ g aliquots were
156	digested with Xyn11 (4 $\mu L$ $\approx$ 21.92 $\mu g;$ Prozomix, UK) for 16 h at 40 $^{\circ}C$ in total volume of
157	$500 \mu$ L. Digestion was terminated by boiling the samples for 30 min and samples were
158	dried in vacuo. Aliquots (200 µg) of undigested XOS/XPS fractions were also dried down.
159	All samples together with standard xylo-oligosaccharides (Xyl <sub>1-6</sub> ; Megazyme, Ireland) were
160	labelled with ANTS and ran on acrylamide gel as described by Kosik, Bromley, Busse-
161	Wicher, Zhang & Dupree (2012). Gels were visualized under UV light using a GelDoc-It
162	TS2 imager (UVP, Germany) equipped with a GFP emission filter (513-557 nm).
163	
164	2.5 In vitro batch fermentation
165	The six different fractions of XOS/XPS along with commercial XOS (avDP2,
166	Shandong Longlive Biotechnology Co. Ltd (SLBC), China) and birch wood xylan (Sigma,
167	Dorset, UK) were evaluated for the ability to modulate the gut microbiota using an <i>in vitro</i>
168	batch culture fermentation system inoculate with human faecal sample.
169	Fructooligosaccharides (avDP 4, Raftilose®, Orafti, Tienan, Belgium) were used as the
170	positive control. The <i>in vitro</i> fermentation was carried out a 50 mL working volume glass
171	jacketed bioreactors, sterile of stirred batch culture fermentation system. The carbohydrates
172	sources were added at 1 % (w/v). The basal medium (per litre) consisted of: 2 g peptone
173	water, 2 g yeast extract, 0.1 g NaCl, 0.04 g K <sub>2</sub> HPO <sub>4</sub> , 0.04 g KH <sub>2</sub> PO <sub>4</sub> , 0.01 g MgSO <sub>4</sub> .7H <sub>2</sub> O,
174	0.01 g CaCl.6H <sub>2</sub> O, 2 g NaHCO <sub>3</sub> , 2 mL Tween 80, 0.05 g haemin, 0.01 mL vitamin K <sub>1</sub> , 0.5
175	g L-cysteine-HCl, 0.5 g bile salt and 4 mL resazurin solution (0.25 g/L).

176	The fermentation of each substrate was carried out in triplicate with each of three
177	healthy human faecal donors, who had not taken prebiotic or probiotic products for 3
178	months, or antibiotics for six months prior to the study. Each vessel containing
179	fermentation medium was inoculated with 5 mL of faecal slurries, which was prior diluted
180	at 10 % (w/w) with anaerobic phosphate-buffered saline (PBS, 0.1 M) and homogenised in
181	a stomacher (Stomacher 400; Seward, West Sussex, UK) for 2 min at medium speed.
182	The fermentation was carried out at pH 6.7-6.9, controlled using an automated pH
183	controller (Fermac 260; Electrolab, Tewkesbury, UK) and at 37 °C (using a
184	thermocirculator) under anaerobic atmosphere, which was achieved through continuous
185	sparging with nitrogen gas. Samples (5 mL) were taken from each fermentation vessel at 0,
186	10, 24 and 36 h for organic acid analysis and bacterial enumeration using the fluorescent in
187	situ hybridisation (FISH) technique.
188	
189	2.6 Enumeration of bacteria
190	The target faecal bacteria groups were enumerated by FISH using 16S rRNA
191	targeted oligonucleotide probes labelled with the fluorescent Cy3 dye. An aliquot (375 $\mu$ L)
192	of sample from each sampling time was mixed with 3 volumes of 4 $\%$ (w/v) cold
193	paraformaldehyde (PFA) solution. The duration of fixation was 5-10 h at 4 $^{\circ}$ C, followed by
194	centrifugation at 13 000 x $g$ for 5 min; the cell pellet was then washed twice with 1 mL cold
195	filter sterilised PBS. The washed cells were then resuspended in 150 $\mu L$ PBS and 150 $\mu L$ of

196 absolute ethanol and stored at -20 °C until analysis.

197	To further process the PFA-fixed sample, 10 $\mu$ L of each sample was diluted with
198	PBS/SDS (sodium dodecyl sulphate) diluent and the diluted samples (20 $\mu$ L) were applied
199	onto six-well of a polytetrafluoroethylene/poly-1-lysine coated slide (Tekdon Inc., Myakka
200	City, FL). The samples were dried at 48 °C for 15 min in a desktop plate incubator and then
201	dehydrated using a series of ethanol solution at 50 %, 80 % and 96 % (v/v) for 3 min each.
202	The excess ethanol was evaporated by drying the slides in a desktop plate incubator for 2
203	min followed by addition of 50 $\mu$ L of mixed hybridisation solution (5 $\mu$ L oligonucleotide
204	probe solution and 45 $\mu$ L hybridisation buffer) onto each well. The slide with samples were
205	hybridised in a microarray hybridisation incubator (Grant-Boekel, Cambridge, UK) for 4 h,
206	washed in 50 mL washing buffer for 15 min and dipped in cold distilled water for 2 s.
207	Slides were dried with compressed air and a drop of PVA-DABCO antifade (polyvinyl
208	alcohol mounting medium with 1, 4-diazabicyclo (2.2.2) octane) was added onto each well.
209	The microscope cover slip was placed on each slide and the cell numbers of
210	microorganisms were determined by direct counting under an epifluorescence microscope
211	(Eclipse 400; Nikon, Surrey, UK) with Fluor 100 lens. A total of 15 fields of view were
212	counted for each well.
213	The probes used were Bif164 (Langendijk et al., 1995), Bac303 (Manz, Amann,
214	Ludwig, Vancanneyt & Schleifer, 1996), Lab158 (Harmsen, Elfferich, Schut & Welling,
215	1999), Ato291 (Harmsen, et al., 2000), Prop853 (Walker, Duncan, McWilliam Leitch,
216	Child & Flint, 2005), Erec482 (Franks et al., 1998), Rrec584 (Walker et al., 2005),
217	Fprau655 (Hold, Schwiertz, Aminov, Blaut & Flint, 2003), Chis150 (Franks et al., 1998),
218	and mixed Eub338 I, II, III (Daims, Brühl, Amann, Schleifer & Wagner, 1999) for
219	enumerating Bifidobacterium spp., Bacteroides-Prevotella, Lactobacillus-Enterococcus,

Atopobium cluster, propionate producing bacteria (Clostridium cluster IX), *Eubacterium rectale-Clostridium cocoides* group (Clostridium cluster XIVa and XIVb), *Roseburia* spp.,
 *Faecalibacterium prausnitzii* cluster *Clostridium histolyticum* group (Clostridium cluster I
 and II) and total bacteria, respectively.

224

225 2.7 Organic acid analysis

226 An aliquot (1 mL) of sample from each sampling time was centrifuged at 13 000 x g 227 for 10 min and the supernatant was stored at -20 °C until analysis. Organic acids analysis 228 was performed using an HPLC (1100 series; Agilent, Winnersh, UK) with refractive index 229 detection. Prior to the analysis, the samples, after thawing, were centrifuged at 13 000 x g 230 for 10 min and the supernatants were filtered through a 0.22 µm filter unit. An ion 231 exclusion column, Rezex ROA-Organic Acid H+ (8%) (Phenomenex, Cheshire, UK) was 232 used for the analysis, using 2.5 mM H<sub>2</sub>SO<sub>4</sub> as eluent. The column was heated at 84 °C and 233 the eluent flow rate was set at 0.5 mL/min. The injection volume used was 20 µL with 40 234 min run time. Organic acids were quantified using standard calibration curves for lactate, 235 acetate, propionate, butyrate and valerate at concentrations of 12.5, 25, 50, 75 and 100 mM. 236 Formate was determined using a formate dehydrogenase-based assay kit (Megazyme, 237 Ireland). 238 239 2.8 Carbohydrate assimilation profile during fermentation 240 The assimilation profile for the nine different carbohydrates substrates used for the 241 batch culture fermentations was determined by High Performance Anion-Exchange

261	3. Results and discussion
260	
259	point. Differences were considered to be significant when $p < 0.05$ .
258	changes for each bacterial group concentration at inoculation and subsequent sampling
257	the different substrates. A paired independent t-test was also used to determine significant
256	differences among the bacterial group populations and organic acid concentrations among
255	analysis of variance (ANOVA) and Tukey's posthoc test was used to determine significant
254	Statistical analysis was performed using SPSS for Windows, version 17.0. One-way
253	2.9 Statistical analysis
252	
251	were used for identification.
250	Standard glucose, arabinose, xylose and xylose oligomers (DP 2-6, Megazyme, Ireland)
249	was then equilibrated for 20 min with 8.75 mM NaOH and 2.4 mM sodium acetate.
248	performed with 125 mM NaOH and 500 mM sodium acetate for 10 min and the column
247	NaOH and 150 mM sodium acetate from 45-49 min. After 50 minutes, a washing step was
246	from 0-19 min; 30 mM NaOH and 25 mM sodium acetate from 20-44 min; and 96.875 mM
245	mL/min with the following linear gradient: 8.75 mM NaOH and 2.4 mM sodium acetate
244	water at a dilution factor of 30. The injection volume was 25 $\mu$ L and the flow rate 1
243	Pulsed Amperometric Detection (PAD). Samples were filtered and diluted with deionised
242	Chromatography (HPAEC, Dionex, Camberley, UK) using a CarboPac PA-1 column and

262 3.1 Characterization of the XOS/XPS fractions

263 The chemical analysis of the six fractions of purified and freeze dried OPEFB
264 autohydrolysis liquor is shown in Table 1. In all cases, XOS/XPS were the dominant
265

#### 266 **Table 1**

267 Composition of OPEFB fibre fractions (g/100 g freeze dried sample) obtained after GFC purification<sup>a</sup>

268

Sample no.	avDP <sup>b</sup>	Residues in linkage (g/100 g)			Ratio <sup>c</sup>		Free Monomers (g/100 g)		Total phenolics (g/100 g)	
	-	Glc	Xyl	Ara	AcO	Ara/Xyl	AcO/Xyl	Xylose	Acetic acid	
1	4	1.75	62.25	1.49	9.16	0.02	0.37	1.23	1.73	0.46
2	7	1.62	65.38	1.16	10.30	0.02	0.39	0.86	0.80	0.37
3	14	1.56	67.32	1.22	11.23	0.02	0.42	0.65	0.48	0.33
4	28	1.61	67.68	1.18	12.43	0.02	0.46	0.57	0.46	0.31
5	44	2.31	64.00	1.21	12.75	0.02	0.50	n.d	0.48	0.43
6	64	2.83	59.28	1.16	12.95	0.02	0.55	n.d	0.46	0.43

<sup>a</sup>In freeze dried form and reconstitute with deionised water to give final concentration of 10 g/L. Calculations were made by assuming the freeze dried samples have 5% moisture content.

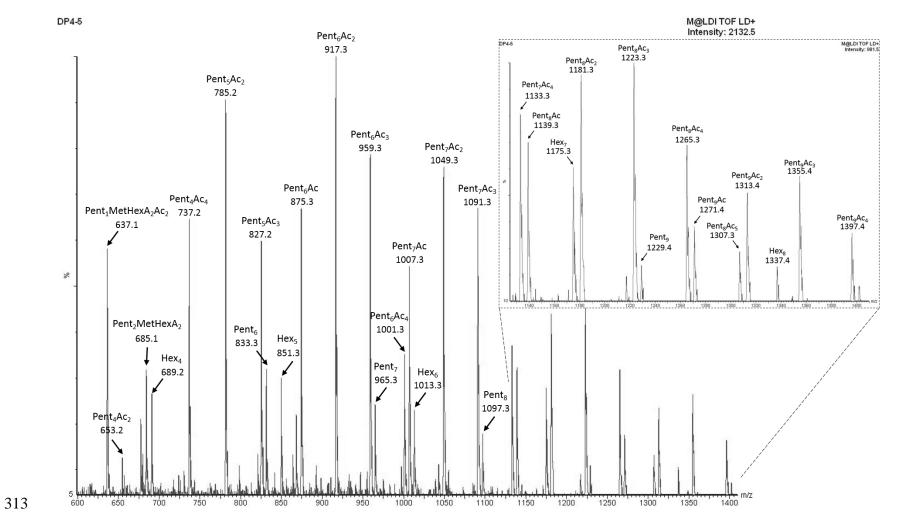
<sup>b</sup>avDP – Average degree of polymerization as determined by size exclusion chromatography

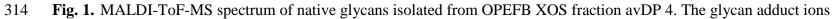
<sup>c</sup>Ratio in mol/mol

AcO - acetyl groups linked to oligosaccharides; n.d. - not detected

270	oligosaccharides, accounting for 78-83 % of the total oligosaccharides. The highest
271	XOS/XPS yield was found in the middle fractions (avDP 14 and 28); free monomeric
272	compounds (xylose and acetic acid) were present at slightly higher concentrations in XOS
273	fractions with lower DP (avDP 4 and 7) than in the other fractions with free xylose absent
274	in XPS fractions with higher DP (avDP 44 and 64). The oligosaccharides of the higher DP
275	fractions (XPS, avDP 44 and 64) were more acetylated. The acetyl groups contribute to the
276	oligosaccharides solubility in water (Nabarlatz, Ebringerová & Montané, 2007) and this
277	may be the reason that high molecular weight XOS fractions were present in OPEFB
278	autohydrolysis liquor. The arabinose content was rather low for all fractions, with an
279	arabinose to xylose ratio of approximating 0.02. The gluco-oligosaccharides (GlcOS) were
280	presumably derived from cellulose and were present at 2-3% w/w. There was also a small
281	amount of total phenolic compounds (<0.5 % w/w) found in all samples.
282	OPEFB fractions (avDP 4, 7 and 14) were analysed by MALDI-ToF-MS (larger
283	avDP fractions were too large for MALDI-ToF-MS analysis). XOS/XPS fractions were all
284	analysed in both their native and permethylated forms by MALDI-ToF-MS. All XOS/XPS
285	fractions analysed in their native form showed acetylated pentose oligosaccharide ions
286	(labelled Pent <sub>n</sub> Ac <sub>n</sub> , the $_{n}$ denoting the number of pentose (Pent) or acetyl (Ac) groups
287	respectively). In avDP 4 the most dominant ion is $m/z$ 917.27 (Pent <sub>6</sub> Ac <sub>2</sub> ) (Fig. 1) with
288	acetylated oligosaccharides ranging from Pent <sub>4</sub> Ac <sub>2</sub> ( $m/z$ 653.19) to Pent <sub>9</sub> Ac <sub>4</sub> ( $m/z$ 1397.42).
289	Also present are pentose oligosaccharides with no acetylation or other modifications with
290	DP 6 to 9 ( $m/z$ 833.25 to 1229.38) and hexose oligosaccharides of DP 4-8 ( $m/z$ 689.21 to
291	1337.42). There could also be small pentose oligosaccharides with methylated-glucuronic
292	acid substitutions (ions at $m/z$ 637.18 and 685.18) found in the native avDP 4 fraction. The

293	permethylated version of avDP 4 fraction (data not shown); although the acetylated
294	residues of the pentose oligosaccharides are lost, we were able to see a pentose ladder
295	starting from Pent <sub>3</sub> ( $m/z$ 549.25) up to Pent <sub>9</sub> ( $m/z$ 1509.69) and ladder of pentose
296	oligosaccharide substituted with single glucuronic acid up to DP 8 (Pen <sub>1</sub> HexA <sub>1</sub> , $m/z$ 447.18
297	to Pen <sub>7</sub> HexA <sub>1</sub> , $m/z$ 1407.63) that could not be observed in native form of the sample.
298	Similarly to the native version of avDP 4 XOS fraction hexose oligosaccharide ladder was
299	observed (Hex <sub>3</sub> $m/z$ 681.33- Hex <sub>8</sub> $m/z$ 1701.83) These data confirm the data in Table 1
300	which showed gluco-oligosaccharides (hexose oligosaccharides), xylo- and arabino-
301	oligosaccharides (pentose oligosaccharides) and acetylated oligosaccharides. Mass
302	spectrometry of OPEFB fractions of avDP 7 and avDP 14 also confirmed the data in Table
303	1. The predominant ions were the acetylated pentoses e.g. $m/z$ 785.18 (Pent <sub>5</sub> Ac <sub>2</sub> ) up to
304	Pent <sub>9</sub> Ac <sub>5</sub> ion ( $m/z$ 1439.43) and methylated glucuronic acid substituted oligosaccharides
305	were also present ( $m/z$ 637.18 and $m/z$ 685.18) (Supplementary Fig. 1a). The permethylated
306	avDP7 fraction (Supplementary Fig. 1b) also contained hexose oligosaccharides (Hex <sub>4</sub> , $m/z$
307	885.43 to Hex <sub>10</sub> $m/z$ 2110.03) as well as pentose oligosaccharide substituted with
308	glucuronic acid (Pent <sub>2</sub> HexA <sub>1</sub> , $m/z$ 607.26 to Pent <sub>9</sub> HexA <sub>1</sub> , $m/z$ 1727.77). In OPEFB the
309	avDP 14 fraction (Supplementary Fig. 1c) acetylated pentose oligosaccharides range from
310	Pent <sub>5</sub> Ac <sub>2</sub> , $m/z$ 785.23 to Pent <sub>18</sub> Ac <sub>6</sub> , $m/z$ 2669.82. Also, observed in the permethylated
311	avDP14 (Supplementary Fig.1d) are glucuronic acid substituted pentoses, Pent <sub>3</sub> HexA <sub>1</sub> ( $m/z$
312	767.33) to Pent <sub>13</sub> HexA <sub>1</sub> ( $m/z$ 2368.07).



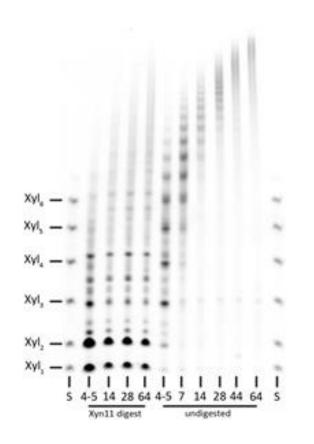


 $[M+Na]^+$  are indicated for acetylated pentose oligosaccharides (Pent<sub>n</sub>Ac<sub>n</sub>), for pentose oligosaccharides (Pent<sub>n</sub>), for pentoses with



The DP ranges of the OPEFB fractions obtained from MALDI-ToF-MS analysis were as follows: avDP 4 (DP 2-9), avDP 7 (DP 3-12), avDP 14 (DP 3-18). The OPEFB fractions were also xylanase cleaved and visualised by polysaccharide analysis using carbohydrate gel electrophoresis (PACE) (Fig.2) which confirms the predominant oligosaccharides were xylo-oligosaccharides and that the gel filtration fractionation of the avDP 4 to avDP 64 contained similar oligosaccharides but with increasing xylose chain length.

324



325

Fig. 2. PACE gel showing separation of extracted OPEFB XOS fractions digested with Xyn11 and undigested. S - Standard xylose<sub>1-6</sub> ladder; 4-5 = avDP 4; 14 = avDP 14; 28 = avDP 28 and 64 = avDP 64, digested with Xyn11. 4-5 = avDP 4; 7 = avDP 7; 14 = avDP 14; 28 = avDP 28; 44 = avDP 44 and 64 = avDP 64, undigested OPEFB XOS fractions.

332	Changes in the bacterial populations during the in vitro fermentations with the
333	different XOS fractions are shown in Table 2. A significant increase (p<0.05) of
334	Bifidobacterium population, ranging between 0.5-0.8 log cells/mL for all time points
335	compared to time 0 h was observed for the XOS fractions with avDP of 4, 7 and 14,
336	commercial XOS and FOS. In the case of the XOS fractions with avDP of 28 and 44,
337	significant increases (p< $0.05$ ) were observed for the 10 h sample, whereas for the XOS
338	fraction with avDP of 64, although an increase was observed for the 10 h sample, this was
339	not statistically significant ( $p \ge 0.05$ ). For all these higher avDP (28, 44, 64) fractions, the
340	concentrations were sustained for the 24 h and 36 h samples and were not statistically
341	different to 0 h. Taking into account the above and the fact that the effect of the XOS
342	fractions with low avDP (avDP 4-14) on the Bifidobacterium population was similar to that
343	of commercial XOS, which mainly consists of DP 2-3, it can be inferred that bifidobacteria
344	preferred the lower molecular weight XOS fractions. This is also supported by the fact that
345	birch wood xylan did not have a significant effect on the Bifidobacterium population. In the
346	pure culture study, there were few strains of Bifidobacterium capable of fermenting high
347	molecular weight XOS or xylan (Palframan, Gibson & Rastall, 2003b). The reason for the
348	increase in the <i>Bifidobacterium</i> population at 10 h for the XOS fractions of avDP 14, 28, 44
349	could be that the bifidobacteria utilise the low molecular weight XOS, which were present
350	in the fractions as demonstrated by the MALDI-ToF-MS. Another possibility is that higher
351	molecular weight XOS was hydrolysed to smaller XOS molecules by other microorganisms
352	such as Bacteroides (Chassard, Goumy, Leclerc, Del'homme & Bernalier-Donadille, 2007;

#### Table 2

Mean bacterial populations in pH-controlled batch cultures at 0, 10, 24 and 36 h<sup>a</sup>

Probe	Time (h)Bacterial population (log10 cells/ml batch culture fluid) in substrate									
		OPEFB XOS	OPEFB XOS	OPEFB XOS	OPEFB XPS	OPEFB XPS	OPEFB XPS	Birch wood	XOS	FOS
		(avDP 4)	(avDP 7)	(avDP 14)	(avDP 28)	(avDP 44)	(avDP 64)	xylan	(SLBC)	(Raftilose)
Bif164	10	8.38 (0.19) <sup>ab*</sup>	8.37 (0.18) <sup>ab*</sup>	8.41 (0.27) <sup>ab*</sup>	8.31 (0.16) <sup>ab*</sup>	8.26 (0.16) <sup>ab*</sup>	8.22 (0.10) <sup>ab</sup>	8.15 (0.10) <sup>a</sup>	8.65 (0.13) <sup>b**</sup>	8.64 (0.08) <sup>b**</sup>
	24	8.56 (0.14) <sup>a*</sup>	8.50 (0.19) <sup>a*</sup>	8.59 (0.16) <sup>a*</sup>	8.40 (0.29) <sup>a</sup>	8.36 (0.28) <sup>a</sup>	8.29 (0.28) <sup>a</sup>	8.25 (0.29) <sup>a</sup>	8.53 (0.06) <sup>a**</sup>	8.48 (0.12) <sup>a*</sup>
7.85(0.09)	36	8.41 (0.15) <sup>a*</sup>	8.46 (0.13) <sup>a*</sup>	8.54 (0.10) <sup>a**</sup>	8.30 (0.24) <sup>a</sup>	8.24 (0.21) <sup>a</sup>	8.10 (0.21) <sup>a</sup>	8.01 (0.23) <sup>a</sup>	8.38 (0.19) <sup>a*</sup>	8.31 (0.35) <sup>a</sup>
Bac303	10	8.58 (0.08) <sup>a*</sup>	8.62 (0.17) <sup>a</sup>	8.64 (0.27) <sup>a</sup>	8.62 (0.13) <sup>a*</sup>	8.46 (0.26) <sup>a</sup>	8.43 (0.14) <sup>a*</sup>	8.48 (0.34) <sup>a</sup>	8.54 (0.11) <sup>a</sup>	8.63 (0.20) <sup>a</sup>
	24	8.50 (0.14) <sup>a**</sup>	8.50 (0.06) <sup>a**</sup>	8.71 (0.04) <sup>a*</sup>	8.59(0.25) <sup>a*</sup>	8.50 (0.44) <sup>a</sup>	8.41 (0.50) <sup>a</sup>	8.59 (0.35) <sup>a</sup>	8.42 (0.13) <sup>a*</sup>	8.46 (0.21) <sup>a</sup>
8.10(0.09)	36	8.30 (0.17) <sup>a</sup>	8.31 (0.12) <sup>a</sup>	8.46 (0.04) <sup>a**</sup>	8.33 (0.32) <sup>a</sup>	8.43 (0.29) <sup>a</sup>	8.27 (0.46) <sup>a</sup>	8.32 (0.14) <sup>a*</sup>	8.29 (0.20) <sup>a</sup>	8.15 (0.08) <sup>a</sup>
Lab158	10	8.30 (0.19) <sup>a</sup>	8.42 (0.25) <sup>a</sup>	8.45 (0.23) <sup>a*</sup>	8.45 (0.20) <sup>a*</sup>	8.38 (0.26) <sup>a</sup>	8.29 (0.05) <sup>a*</sup>	8.27 (0.14) <sup>a</sup>	8.51 (0.13) <sup>a*</sup>	8.45 (0.22) <sup>a</sup>
	24	8.36 (0.17) <sup>a</sup>	8.50 (0.19) <sup>a*</sup>	8.57 (0.20) <sup>a*</sup>	8.46 (0.07) <sup>a**</sup>	8.29 (0.35) <sup>a</sup>	8.24 (0.24) <sup>a</sup>	8.42 (0.19) <sup>a*</sup>	8.35 (0.14) <sup>a*</sup>	8.30 (0.15) <sup>a*</sup>
7.97(0.04)	36	8.31 (0.17) <sup>a</sup>	8.45 (0.12) <sup>a*</sup>	8.46 (0.14) <sup>a*</sup>	8.27 (0.11) <sup>a</sup>	8.10 (0.31) <sup>a</sup>	8.04 (0.32) <sup>a</sup>	8.13 (0.19) <sup>a</sup>	8.28 (0.24) <sup>a</sup>	8.32 (0.35) <sup>a</sup>
Ato291	10	8.22 (0.03) <sup>ab*</sup>	8.19 (0.05) <sup>ab*</sup>	8.12(0.05) <sup>ab**</sup>	8.07 (0.18) <sup>a</sup>	8.00 (0.07) <sup>a</sup>	8.05 (0.14) <sup>a</sup>	7.97 (0.32) <sup>a</sup>	8.42 (0.17) <sup>ab**</sup>	8.56 (0.20) <sup>b**</sup>
	24	8.14 (0.09) <sup>bcd*</sup>	8.08 (0.10) <sup>abc**</sup>	7.93 (0.11) <sup>ab</sup>	7.99 (0.03) <sup>abc*</sup>	7.87 (0.04) <sup>ab</sup>	7.72 (0.05) <sup>a</sup>	8.00 (0.20) <sup>abc</sup>	8.35 (0.20) <sup>cd*</sup>	8.51 (0.23) <sup>d**</sup>
7.78(0.10)	36	7.88 (0.23) <sup>abc</sup>	7.80 (0.17) <sup>ab</sup>	7.81 (0.16) <sup>ab</sup>	7.69 (0.10) <sup>a</sup>	7.60 (0.15) <sup>a</sup>	7.57 (0.15) <sup>a</sup>	7.66 (0.21) <sup>a</sup>	8.22 (0.24) <sup>bc</sup>	8.37 (0.05) <sup>c**</sup>
Prop853	10	7.90 (0.04) <sup>a*</sup>	8.07 (0.12) <sup>a</sup>	$8.08(0.08)^{a^*}$	8.11 (0.05) <sup>a**</sup>	8.05 (0.03) <sup>a**</sup>	8.01 (0.23) <sup>a</sup>	7.92 (0.32) <sup>a</sup>	7.99 (0.23) <sup>a</sup>	7.97 (0.25) <sup>a</sup>
-	24	8.03 (0.26) <sup>a</sup>	8.12 (0.09) <sup>a*</sup>	8.17 (0.08) <sup>a*</sup>	8.13 (0.14) <sup>a*</sup>	8.04 (0.30) <sup>a</sup>	7.87 (0.37) <sup>a</sup>	7.98 (0.23) <sup>a</sup>	8.02(0.32) <sup>a</sup>	7.97 (0.37) <sup>a</sup>
7.71(0.05)	36	7.87 (0.33) <sup>a</sup>	7.92 (0.16) <sup>a</sup>	7.86 (0.19) <sup>a</sup>	7.78 (0.13) <sup>a</sup>	7.74 (0.41) <sup>a</sup>	7.68 (0.41) <sup>a</sup>	7.76 (0.20) <sup>a</sup>	7.61 (0.12) <sup>a</sup>	7.86 (0.24) <sup>a</sup>
Erec482	10	8.09 (0.20) <sup>a</sup>	8.18 (0.47) <sup>a</sup>	8.28 (0.48) <sup>a</sup>	8.29 (0.37) <sup>a</sup>	8.15 (0.18) <sup>a</sup>	8.20 (0.14) <sup>a</sup>	8.20 (0.18) <sup>a</sup>	8.28 (0.30) <sup>a</sup>	8.31 (0.24) <sup>a</sup>
	24	8.26 (0.12) <sup>a*</sup>	8.44 (0.34) <sup>a</sup>	8.35 (0.51) <sup>a</sup>	8.43(0.27) <sup>a</sup>	8.08 (0.52) <sup>a</sup>	8.22 (0.27) <sup>a</sup>	8.24 (0.14) <sup>a*</sup>	8.36 (0.15) <sup>a*</sup>	8.33 (0.08) <sup>a**</sup>
7.99(0.04)	36	8.43 (0.10) <sup>a*</sup>	8.41 (0.33) <sup>a</sup>	8.27 (0.47) <sup>a</sup>	8.28 (0.09) <sup>a*</sup>	8.19 (0.32) <sup>a</sup>	8.13 (0.41) <sup>a</sup>	8.20 (0.33) <sup>a</sup>	8.28 (0.14) <sup>a*</sup>	8.14 (0.12) <sup>a</sup>
Rrec584	10	7.48 (0.16) <sup>a</sup>	7.48 (0.11) <sup>a</sup>	7.49 (0.18) <sup>a</sup>	7.45 (0.02) <sup>a*</sup>	7.38 (0.02) <sup>a*</sup>	7.35(0.06) <sup>a*</sup>	7.38 (0.12) <sup>a</sup>	7.52(0.22) <sup>a</sup>	7.41 (0.17) <sup>a</sup>
	24	7.61 (0.06) <sup>ab</sup>	7.58 (0.11) <sup>ab</sup>	7.46 (0.19) <sup>a</sup>	7.54 (0.10) <sup>ab</sup>	7.51 (0.17) <sup>ab</sup>	7.50(0.06) <sup>a</sup>	7.50(0.15) <sup>a</sup>	7.85 (0.05) <sup>b*</sup>	$7.76(0.11)^{ab^*}$
7.38(0.05)	36	7.70 (0.22) <sup>a*</sup>	7.65 (0.15) <sup>a</sup>	7.65 (0.21) <sup>a</sup>	7.53 (0.07) <sup>a*</sup>	7.59 (0.20) <sup>a</sup>	7.60 (0.12) <sup>a</sup>	7.40 (0.20) <sup>a</sup>	7.87 (0.20) <sup>a*</sup>	7.75 (0.15) <sup>a</sup>
Fprau655	10	7.58 (0.26) <sup>a</sup>	7.67 (0.29) <sup>a</sup>	7.66 (0.30) <sup>a</sup>	7.72 (0.13) <sup>a</sup>	7.61 (0.19) <sup>a</sup>	7.62 (0.13) <sup>a</sup>	7.65 (0.30) <sup>a</sup>	7.53 (0.26) <sup>a</sup>	7.67 (0.34) <sup>a</sup>
	24	7.36 (0.08) <sup>a</sup>	7.45 (0.11) <sup>a</sup>	7.57 (0.24) <sup>a</sup>	7.84 (0.10) <sup>a*</sup>	7.51 (0.27) <sup>a</sup>	7.58 (0.22) <sup>a</sup>	7.74 (0.27) <sup>a</sup>	7.49 (0.20) <sup>a</sup>	7.60 (0.22) <sup>a</sup>
7.54(0.10)	36	7.44 (0.24) <sup>a</sup>	7.46 (0.02) <sup>a</sup>	7.47 (0.21) <sup>a</sup>	7.56 (0.19) <sup>a</sup>	7.40 (0.11) <sup>a*</sup>	7.55 (0.27) <sup>a</sup>	7.48 (0.28) <sup>a</sup>	7.34 (0.12) <sup>a</sup>	7.43 (0.25) <sup>a</sup>
Chis150	10	7.41 (0.21) <sup>a</sup>	7.48 (0.09) <sup>a</sup>	7.44 (0.13) <sup>a</sup>	7.49 (0.10) <sup>a</sup>	7.38 (0.07) <sup>a</sup>	7.38 (0.15 ) <sup>a</sup>	7.44 (0.09) <sup>a</sup>	7.36 (0.11) <sup>a</sup>	7.56 (0.27) <sup>a</sup>
	24	7.34 (0.04) <sup>a</sup>	7.34 (0.15) <sup>a</sup>	7.23 (0.06) <sup>a</sup>	7.31 (0.10) <sup>a</sup>	7.27 (0.05) <sup>a</sup>	7.24 (0.07) <sup>a</sup>	7.36 (0.09) <sup>a</sup>	7.28 (0.03) <sup>a</sup>	7.34 (0.12) <sup>a</sup>
7.33(0.05)	36	6.93 (0.12) <sup>a</sup>	6.97 (0.08) <sup>a*</sup>	6.95 (0.15) <sup>a</sup>	6.91 (0.06 ) <sup>a*</sup>	6.90 (0.16) <sup>a</sup>	6.71 (0.07) <sup>a*</sup>	6.95 (0.15) <sup>a*</sup>	6.88 (0.08) <sup>a*</sup>	6.80 (0.06 ) <sup>a*</sup>
Eub338	10	9.17 (0.11) <sup>a</sup>	9.21 (0.12) <sup>a*</sup>	9.22 (0.10) <sup>a*</sup>	9.19 (0.06) <sup>a*</sup>	9.16 (0.15) <sup>a</sup>	9.10 (0.10) <sup>a*</sup>	9.06 (0.14) <sup>a</sup>	9.33 (0.18) <sup>a</sup>	9.30 (0.11) <sup>a*</sup>
	24	9.25(0.17) <sup>a</sup>	9.25 (0.10) <sup>a*</sup>	9.30 (0.14) <sup>a*</sup>	9.25 (0.13) <sup>a*</sup>	9.24 (0.18) <sup>a</sup>	9.12 (0.13) <sup>a</sup>	9.12 (0.16) <sup>a</sup>	9.27 (0.16) <sup>a</sup>	9.18 (0.11) <sup>a</sup>
8.79(0.06)	36	9.16(0.19) <sup>a</sup>	9.15(0.15) <sup>a</sup>	9.22 (0.13) <sup>a</sup>	9.08 (0.05) <sup>a*</sup>	9.08 (0.34) <sup>a</sup>	8.93 (0.33) <sup>a</sup>	8.91 (0.19) <sup>a</sup>	9.09 (0.17) <sup>a</sup>	9.06 (0.13) <sup>a</sup>

<sup>a</sup>Standard deviation is shown in parentheses (n=3). Significant differences (p<0.05) between substrates are indicated with different letters in a same row. <sup>\*</sup>Significant differences from value at 0 h, p<0.05; <sup>\*\*</sup>Significant differences from value at 0 h, p<0.01 (Value at 0 h is shown in the far left under 'Probe' column)

354 Falony, Calmeyn, Leroy & De Vuyst, 2009). This was also observed in studies carried out 355 by Mäkeläinen and co-workers (2010a; 2010b), a high molecular weight xylan (DP 35-40) 356 was not efficiently metabolised by a range of *Bifidobacterium* strains in pure culture studies 357 but when they tested the same xylan in a semi continuous colon simulator system using 358 faecal inoculum, they observed a significant increase in the *Bifidobacterium* sp. population. 359 Another bacterial group which had significant difference between substrates is the 360 Atopobium cluster. Atopobium has the highest count on FOS, significantly higher (p<0.05) 361 than OPEFB XOS of avDP 28, 44 and 64. These results are consistent with Hughes et al. 362 (2007) whereby the large molecular weight AXOS (278 kDa and 354 kDa) generally did 363 not induce growth of Atopobium.

364

365 3.3 Organic acid analysis

Table 3 shows the organic acid concentrations in the fermentations; acetate was the leading SCFA produced, followed by propionate, formate, lactate and butyrate. Across all substrates, formate and lactate were transient metabolites reaching maximum at 10 h. Acetate and propionate concentration on the other hand continued to rise up to 24 h and/or 36 h, whereas butyrate, though present at low concentration initially, increased steadily up to 36 h.

All OPEFB XOS produced significantly lower (p<0.05) amount of lactate than commercial XOS and FOS. The wider DP distribution and possibility the presence of substituents on OPEFB XOS may affect the accessibility for bifidobacterial fermentation. Kabel et al. (2002a) also observed a higher amount of lactate in non-substituted XOS than 376 substituted XOS. According to Falony et al. (2009), metabolism in bifidobacteria produces 377 more formate, acetate and ethanol at the expense of lactate when there is limited access to 378 substrate. Different carbohydrates are known to promote the growth of different species of 379 bifidobacteria, resulting in varying amount of lactate (Palframan et al., 2003b). 380 The initial acetate level in OPEFB XOS avDP 4 was high, possibly as a result of 381 free acetic acid present in the low molecular weight substrate. XOS in all OPEFB fractions 382 and the commercial XOS resulted in higher acetate and less propionate and butyrate than 383 FOS. This typical profile corresponds with previous studies conducted on XOS and xylan

- fermentation (Englyst, Hay & Macfarlane, 1987; Kabel et al., 2002a; Rycroft, Jones,
- 385 Gibson & Rastall, 2001).

#### 386 Table 3 Mean organic acid concentrations in pH-controlled batch cultures at 0, 10, 24 and 36 h<sup>a</sup>

Organic acid	Time				С	oncentration (mM	)			
	(h)	OPEFB XOS	OPEFB XOS	OPEFB XOS	OPEFB XPS	OPEFB XPS	OPEFB XPS	Birch wood	XOS (Suntory)	FOS (Raftilose)
		(avDP 4)	(avDP 7)	(avDP 14)	(avDP 28)	(avDP 44)	(avDP 64)	xylan		× /
Lactate	0	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.78 (0.68) <sup>a</sup>	0.79 (0.68) <sup>a</sup>
	10	4.88 (2.92) <sup>a</sup>	1.85 (2.32) <sup>a</sup>	2.46 (2.38) <sup>a</sup>	0.81 (1.40) <sup>a</sup>	2.34 (1.44) <sup>a</sup>	$0.46 (0.79)^{a}$	0.79 (0.72) <sup>a</sup>	16.11 (5.89) <sup>b*</sup>	19.29 (6.34) <sup>b*</sup>
	24	0.56 (0.98) <sup>a</sup>	0.32 (0.56) <sup>a</sup>	1.02 (1.76) <sup>a</sup>	$0.00 (0.00)^{a}$	0.50 (0.87) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.32 (0.56) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>
	36	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.45 (0.78) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.47 (0.81) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>
Formate	0	0.58 (0.04) <sup>d</sup>	0.16 (0.07) <sup>bc</sup>	0.11 (0.07) <sup>abc</sup>	0.11 (0.06) <sup>abc</sup>	0.13 (0.06) <sup>abc</sup>	0.36 (0.04) <sup>a</sup>	0.16 (0.03) <sup>c</sup>	0.03 (0.04) <sup>ab</sup>	0.01 (0.01) <sup>a</sup>
	10	8.42 (8.28) <sup>a</sup>	7.61 (7.02) <sup>a</sup>	4.26 (5.39) <sup>a</sup>	8.37 (6.06) <sup>a</sup>	4.11 (5.44) <sup>a</sup>	5.80 (7.78) <sup>a</sup>	2.64 (1.64) <sup>a</sup>	14.06 (3.49) <sup>a*</sup>	14.96(5.90) <sup>a*</sup>
	24	5.33(3.65) <sup>a</sup>	5.66 (8.32) <sup>a</sup>	3.54(6.12) <sup>a</sup>	4.26 (7.38) <sup>a</sup>	6.44 (5.58) <sup>a</sup>	2.19 (3.56) <sup>a</sup>	0.05 (0.08) <sup>a</sup>	6.56 (5.94) <sup>a</sup>	1.69 (2.86) <sup>a</sup>
	36	0.00 (0.00) <sup>a</sup>	1.93 (3.34) <sup>a</sup>	0.55 (0.95) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	2.34 (2.54) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	1.24 (2.15) <sup>a</sup>	0.00 (0.00) <sup>a</sup>
Acetate	0	10.08 (2.41) <sup>c</sup>	6.70 (1.79) <sup>bc</sup>	6.00 (1.56) <sup>b</sup>	5.62 (1.72) <sup>ab</sup>	5.20 (0.50) <sup>ab</sup>	5.18 (0.38) <sup>ab</sup>	5.84 (0.06) <sup>ab</sup>	2.18 (0.03) <sup>a</sup>	2.31 (0.30) <sup>a</sup>
(A)	10	48.44 (21.23) <sup>a</sup>	47.45 (24.27) <sup>a</sup>	47.84 (23.12) <sup>a</sup>	51.72 (24.48) <sup>a</sup>	33.12 (22.07) <sup>a</sup>	37.19 (27.31) <sup>a</sup>	25.53 (6.71) <sup>a*</sup>	54.82 (8.47) <sup>a**</sup>	47.55 (11.02) <sup>a*</sup>
	24	77.39 (21.26) <sup>b*</sup>	71.61 (7.48) <sup>ab**</sup>	78.37 (6.57) <sup>b**</sup>	62.35 (11.82) <sup>ab*</sup>	57.30 (28.36) <sup>ab</sup>	43.50 (26.36) <sup>ab</sup>	28.98 (7.96) <sup>a*</sup>	60.19 (2.00) <sup>ab**</sup>	43.10 (6.47) <sup>ab**</sup>
	36	79.80 (22.19) <sup>b*</sup>	68.68 (10.00) <sup>ab**</sup>	78.70 (6.86) <sup>b**</sup>	54.60 (10.09) <sup>ab*</sup>	59.49 (27.07) <sup>ab</sup>	41.44 (30.12) <sup>ab</sup>	21.32 (7.03) <sup>a</sup>	60.53 (3.77) <sup>ab**</sup>	39.61 (7.66) <sup>ab*</sup>
Propionate	0	3.08 (0.52) <sup>a</sup>	2.75 (0.19) <sup>a</sup>	2.69 (0.18) <sup>a</sup>	2.67 (0.19) <sup>a</sup>	2.70 (0.10) <sup>a</sup>	2.69 (0.22) <sup>a</sup>	2.76 (0.18) <sup>a</sup>	2.61 (0.11) <sup>a</sup>	2.67 (0.42) <sup>a</sup>
(P)	10	9.23 (4.64) <sup>a</sup>	13.84 (10.69) <sup>a</sup>	15.60 (11.96) <sup>a</sup>	12.77 (1.82) <sup>a*</sup>	7.96 (1.94) <sup>a*</sup>	11.37 (6.05) <sup>a</sup>	10.28 (2.78) <sup>a*</sup>	13.28 (8.04) <sup>a</sup>	15.55 (14.71) <sup>a</sup>
	24	16.57 (4.75) <sup>a*</sup>	20.10 (7.20) <sup>a</sup>	25.10 (8.72) <sup>a*</sup>	18.43 (2.87) <sup>a*</sup>	18.15 (10.37) <sup>a</sup>	11.46 (10.51) <sup>a</sup>	13.22 (4.72) <sup>a</sup>	18.07 (8.97) <sup>a</sup>	18.58 (16.11) <sup>a</sup>
	36	17.93 (5.55) <sup>a*</sup>	18.82 (6.41) <sup>a*</sup>	25.70 (7.51) <sup>a*</sup>	16.35 (2.54) <sup>a*</sup>	19.27 (11.63) <sup>a</sup>	11.28 (11.08) <sup>a</sup>	9.91 (3.20) <sup>a</sup>	17.96 (9.82) <sup>a</sup>	18.22 (16.91) <sup>a</sup>
Butyrate	0	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>	0.00 (0.00) <sup>a</sup>
(B)	10	1.11 (1.72) <sup>a</sup>	1.85 (1.84) <sup>a</sup>	1.87 (2.33) <sup>a</sup>	2.08 (1.47) <sup>a</sup>	1.11 (1.44) <sup>a</sup>	1.89 (1.06) <sup>a</sup>	1.76 (0.86) <sup>a</sup>	1.89 (1.65) <sup>a</sup>	2.68 (1.08) <sup>a*</sup>
	24	2.99 (1.81) <sup>a</sup>	3.08 (2.13) <sup>a</sup>	3.32 (3.40) <sup>ab</sup>	3.67 (1.74) <sup>ab</sup>	1.66 (1.46) <sup>a</sup>	2.66 (2.48) <sup>a</sup>	3.39 (1.92) <sup>ab</sup>	11.41 (5.31) <sup>bc</sup>	13.16 (3.29) <sup>c*</sup>
	36	4.07 (1.75) <sup>ab</sup>	3.52 (2.31) <sup>a</sup>	4.49 (4.11) <sup>ab</sup>	4.09 (1.03) <sup>ab*</sup>	2.40 (2.02) <sup>a</sup>	3.24 (3.60) <sup>a</sup>	3.09 (2.79) <sup>a</sup>	12.30 (4.64) <sup>bc*</sup>	13.23 (2.49) <sup>c*</sup>
Total	0	13.73 (2.70) <sup>b</sup>	9.61 (1.85) <sup>ab</sup>	8.80 (1.65) <sup>a</sup>	8.40 (1.87) <sup>a</sup>	8.04 (0.42) <sup>a</sup>	8.23 (0.56) <sup>a</sup>	8.77 (0.22) <sup>a</sup>	5.59 (0.67) <sup>a</sup>	5.77 (1.38) <sup>a</sup>
	10	72.09(29.09) <sup>a</sup>	72.60 (33.75) <sup>a</sup>	72.03 (32.92) <sup>a</sup>	75.74 (32.56) <sup>a</sup>	48.64 (27.61) <sup>a</sup>	56.71 (41.11) <sup>a</sup>	41.00(11.33) <sup>a*</sup>	100.15 (7.69) <sup>a**</sup>	100.03 (1.46) <sup>a**</sup>
	24	102.84 (27.40) <sup>a*</sup>	100.77 (7.09) <sup>a**</sup>	111.35 (6.55) <sup>a**</sup>	88.71 (21.28) <sup>a*</sup>	84.05 (44.15) <sup>a</sup>	59.80 (42.27) <sup>a</sup>	45.97(13.62) <sup>a*</sup>	96.24 (4.10) <sup>a**</sup>	76.52 (10.92) <sup>a**</sup>
	36	101.80(26.14) <sup>ab*</sup>	92.95 (6.64) <sup>ab**</sup>	109.44 (2.12) <sup>b**</sup>	75.04 (12.69) <sup>ab*</sup>	83.96 (41.10) <sup>ab</sup>	55.95 (44.40) <sup>ab</sup>	34.79 (12.76) <sup>a</sup>	92.03 (7.57) <sup>ab**</sup>	71.05 (17.60) <sup>ab*</sup>
A:P:B	0	1:0.3:0	1:0.4:0	1:0.5:0	1:0.5:0	1:0.5:0	1:0.5:0	1:0.5:0	1:1.2:0	1:1.2:0
	10	1:0.2:0.03	1:0.3:0.03	1:0.3:0.03	1:0.3:0.04	1:0.3:0.04	1:0.3:0.06	1:0.4:0.07	1:0.3:0.03	1:0.4:0.06
	24	1:0.2:0.04	1:0.3:0.04	1:0.3:0.04	1:0.3:0.06	1:0.3:0.04	1:0.3:0.06	1:0.5:0.1	1:0.3:0.2	1:0.5:0.3
	36	1:0.2:0.05	1:0.3:0.05	1:0.3:0.06	1:0.3:0.08	1:0.3:0.05	1:0.3:0.06	1:0.5:0.1	1:0.3:0.2	1:0.5:0.4

<sup>a</sup>Standard deviation is shown in parentheses with n=3. Significant differences (p<0.05) between substrates are indicated with different letters in a same row. \*Increased significantly from value at 0 h, p<0.05; \*\*Increased significant differences from value at 0 h, p<0.01

387 The significant increase in acetate at 24 h and 36 h for XOS of avDP 4, 7, 14 and 28
388 can be linked to the two major acetate producers; *Bifidobacterium* spp. and the *Bacteroides*389 *Prevotella* group.

390 There was no significant increase in butyrate on any OPEFB XOS while 391 commercial XOS resulted in similar butyrate level to FOS ( $p \ge 0.05$ ). Nevertheless, the XOS 392 preparation of lower avDP (4, 14, 28) were not different to commercial XOS ( $p \ge 0.05$ ). 393 Although the human gut microbiota has also been known to be able to further metabolise 394 acetate to butyrate (Duncan, Barcenilla, Stewart, Pryde & Flint, 2002; Duncan et al., 2004), 395 the conversion of acetate from OPEFB XOS to butyrate was generally lower. 396 The type and molecular weight of the substrates influenced rate and amount of 397 organic acid produced. Based on total organic acid, it is noticeable that commercial XOS 398 and FOS were the fastest fermentable substrates, reaching at least 100 mM at 10 h. As for 399 OPEFB XOS, the three lowest avDP (4, 7, 14) reached 100 mM at 24 h while other 400 fractions of higher avDP (28, 44, 64) had less than 100 mM and birch wood xylan, the least 401 fermentable substrate had the lowest organic acid of all with 46 mM at 24 h. 402 403 3.4 Carbohydrate assimilation profile during fermentation

404 The carbohydrate was profiled in the samples during the course of fermentation using

405 HPAEC-PAD to observe the changes in DP. The assimilation profile of OPEFB XOS of

406 avDP14 from each faecal donor is illustrated in Fig. 3. The three donors showed slight

407 variation in magnitudes and trends that coincides with rather high standard deviation

408 observed in the organic acid data. At 10 h, donor 1 XOS were utilised faster, leaving behind

409	xylose. For donor 2, since the rate of fermentation is slower, some oligosaccharides were
410	still present at 10 h and without much increase of xylose. Donor 3 had a trend between
411	donor 1 and 2 whereby the XOS were also quickly fermented and broken down into xylose,
412	xylobiose and xylotriose. At 24 h there was no detectable sugar remaining in all the culture
413	samples. While the xylose and low DP XOS were being consumed by the bacteria,
414	accumulation could arise from continual digestion of XOS/XPS from the higher DP. This
415	similar degradation characteristic was also observed in XOS (DP 2-6) derived from rice
416	husk when fermented with a single bifidobacteria culture (Gullón et al., 2008).
417	Analysis with HPAEC-PAD however does not provide information on acetyl groups
418	as deacetylation occurs in the high pH eluent used in HPAEC (Kabel et al., 2002a). As such,
419	the chromatogram could not show the susceptibility of acetylated XOS during fermentation.
420	

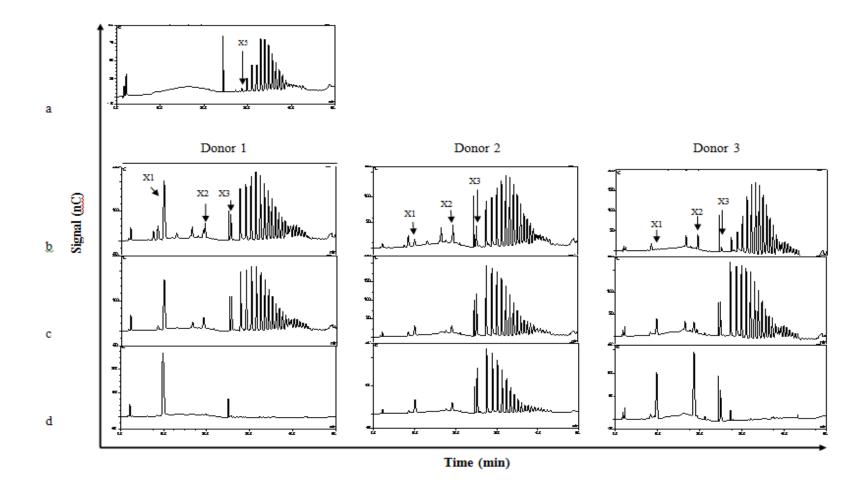


Fig. 3. Degradation profile of OPEFB XOS avDP 14 at different time by faecal culture from three donors using HPAEC-PAD:
(a) Substrate before fermentation, (b) Immediately after substrate addition into fermenter, (c) After 5 h, (d) After 10 h. X1, X2,
X3 on the chromatogram indicate the position of xylose, xylobiose and xylotriose, respectively.

#### **4. Conclusion**

426	The solubility of high avDP XOS/XPS preparation from OPEFB through
427	autohydrolysis process is rather interesting as it could be incorporated into many food
428	processes. The acetyl group may aid XOS/XPS solubility, however the impact of this on
429	fermentation in the gut was not conclusive from the present results. Nevertheless, the
430	degree of polymerisation has significant influence on OPEFB XOS/XPS fermentability by
431	the gut microflora. The <i>in vitro</i> study conducted in this work shows the low avDP XOS (4,
432	7, 14) were more selective to beneficial bacteria than the higher avDP XPS (22, 44, 64).
433	OPEFB XOS fractions of avDP 14 appeared to be the most bifidogenic.
434	
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437 438 439	scholarship to Ai Ling Ho. The XOS fractions used in this study was produced using the facilities provided at Bioenergy Unit, LNEG, Portugal. Rothamsted Research receives grant-aided support from the Biotechnology and Biological Sciences Research Council
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437 438 439 440 441	scholarship to Ai Ling Ho. The XOS fractions used in this study was produced using the facilities provided at Bioenergy Unit, LNEG, Portugal. Rothamsted Research receives grant-aided support from the Biotechnology and Biological Sciences Research Council (BBSRC) of the UK.
<ul> <li>437</li> <li>438</li> <li>439</li> <li>440</li> <li>441</li> <li>442</li> </ul>	scholarship to Ai Ling Ho. The XOS fractions used in this study was produced using the facilities provided at Bioenergy Unit, LNEG, Portugal. Rothamsted Research receives grant-aided support from the Biotechnology and Biological Sciences Research Council (BBSRC) of the UK. References

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