



**STUDY OF THE EFFECT OF DIETARY FIBER FRACTIONS
OBTAINED FROM ARTICHOKE (*Cynara cardunculus* L. var.
scolymus) ON THE GROWTH OF INTESTINAL BACTERIA
ASSOCIATED WITH HEALTH**

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23 **ABSTRACT**

24

25 The effect of different fractions enriched in soluble fiber obtained from
26 artichoke using citric acid or citric acid / hemicellulase, on the selective growth of
27 *Lactobacillus plantarum* 8114 and *Bifidobacterium bifidum* ATCC 11863 was
28 evaluated. Gompertz modeling of *Lactobacillus plantarum* 8114 growth showed a
29 higher specific growth rate (μ : 0.16 h⁻¹) in the presence of fraction isolated from stem
30 using hemicellulase (fraction A) than in the presence of glucose (μ : 0.09 h⁻¹). In the
31 case of *Bifidobacterium bifidum* 11863, the highest μ was obtained for the
32 microorganism grown in the presence of fraction A and for the fraction isolated from
33 stem without hemicellulase, being their rate twice the one observed for glucose (0.04
34 h⁻¹). The positive prebiotic activity scores observed with respect to *Escherichia coli*
35 25922 indicated that fibers assayed are metabolized as well as glucose by
36 *Lactobacillus plantarum* 8114 and *Bifidobacterium bifidum* ATCC 11863 and that
37 they are selectively metabolized by these microorganisms. The potential capacity for
38 selectively stimulate the growth of intestinal bacteria associated with health shown by
39 fraction A can be ascribed to its high inulin and low methylation degree pectin
40 contents.

41

42

43 **Keywords:** artichoke; lactobacilli and bifidobacteria; prebiotic activity score.

44

45 1. Introduction

46

47 The human gut microflora is affected by many factors such as age, drug
48 therapy, diet, host physiology, peristalsis, local immunity and “in situ” bacterial
49 metabolism.¹ However; diet is probably the most significant factor determining the
50 type of gut flora that develops since foodstuffs provide the main nutrient sources for
51 colonic bacteria. This has led to the concept of prebiotics. A prebiotic was first
52 defined as a ‘non-digestible food ingredient that beneficially affects the host by
53 selectively stimulating the growth and/or activity of one or a limited number of
54 bacteria in the colon, and thus improves host health’.² In particular, many food
55 oligosaccharides and polysaccharides have been claimed to have prebiotic activity,
56 but not all dietary carbohydrates are prebiotics.³

57 Roberfroid⁴ stated that the classification of a food ingredient as a prebiotic
58 requires a scientific demonstration that the ingredient:

59 (1) resists gastric acidity, hydrolysis by mammalian enzymes and gastrointestinal
60 absorption;

61 (2) is fermented by the intestinal microflora;

62 (3) stimulates selectively the growth and/or activity of intestinal bacteria associated
63 with health and wellbeing.

64 As the field of prebiotics has developed, so has the methodology for
65 investigating functionality. In general, the changes of flora in response to diet have
66 been studied using strains of *Bifidobacterium* spp. and *Lactobacillus* spp. and
67 comparing its growth with the one of other bacteria such as *Bacteroides* spp.,
68 *Clostridium* spp., *Eubacterium* spp. and *Escherichia coli*.⁵ The number of strains
69 tested varies with different reports. Currently, it is proposed to evaluate the fulfillment
70 of the three requirements previously mentioned for defining a food ingredient as a
71 prebiotic, being the selective stimulation of growth the first stage in the evaluation of
72 the characteristics of different food ingredients.³ For example, Marotti et al.⁶ studied
73 the prebiotic effect of soluble fibers from seven modern, two old and one ancient
74 durum-type wheat varieties on *Lactobacillus* and *Bifidobacterium* strains. In that
75 study, the behaviors of *L. plantarum* L12 and *B. pseudocatenulatum* B7003 were
76 studied in the presence of wheat fiber and glucose and compared with the behavior of
77 *Escherichia coli* ATCC 25645 and *Klebsiella pneumoniae* GC 23a in the presence of
78 both carbon sources to evaluate the prebiotic activity of wheat fiber. Fiber microbial

79 utilization was highly variable and dependent on the strain. Soluble dietary fibers
80 from durum-type wheat grains were identified as potential prebiotic substrate for the
81 selective proliferation of *B. pseudocatenulatum* B7003 and *L. plantarum* L12 *in vitro*.
82 Several studies have shown that the ability of lactobacilli and bifidobacteria to
83 ferment prebiotic carbohydrates is both strain and substrate specific.^{7,8} In addition, it
84 is not clear which prebiotic carbohydrates are the most suitable substrates for selective
85 growth of specific strains. Recently, several quantitative approaches were devised to
86 determine the functional activity of prebiotics during *in vitro* fermentation conditions.
87 In general, these studies provided indices that reflect the relative ability of a given
88 prebiotic to produce specific effects, and are based on the measurement of microbial
89 populations, growth rates, substrate assimilation, and/or short-chain fatty acid
90 production. The indices were then used to rank various carbohydrates according to
91 their potential to stimulate growth of specific members of a mixed microflora.
92 However, as fermentation of prebiotics is dependent on the bacterial strain, rather than
93 based on the species or genera, it is important to understand the extent to which the
94 metabolism of prebiotics occurs by specific strains of bacteria, especially for those
95 organisms whose intended use is as probiotics.^{9,10,11,12}

96 In a previous work, Fissore et al.¹³ reported the antioxidant and *in vitro*
97 antiviral effects of dietary fiber fractions isolated with citric acid or citric acid /
98 hemicellulase from bracts, stems and hearts of artichoke (*Cynara cardunculus* L. var.
99 *scolymus*). These fractions contained inulin and pectin. The aim of the present study is
100 to quantify the extent to which those fractions selectively stimulate the growth of the
101 strains *Lactobacillus plantarum* 8114 and *Bifidobacterium bifidum* ATCC 11863 with
102 the purpose of helping in the understanding of the potential of different fibers to act as
103 prebiotic substrates. In addition, the kinetic parameters of microbial growth were also
104 studied.

105

106 **2. Materials and Methods**

107 *2.1. Sample preparation*

108 Artichokes (*Cynara cardunculus* L. var. *scolymus*) harvested in Argentina
109 were bought in the local market. Bracts, hearts and stems were separated, washed with
110 distilled water, dried (85 °C, 2.5 h) in a convection oven (0.508 m/s of air rate), milled
111 (E909, Wemir, Buenos Aires, Argentina) and sieved for obtaining powders enriched
112 in cell wall material (CWM) with particle sizes in the range 420 - 710 µm.

113 Each CWM was treated as follows:

114 10 g of CWM were poured into a beaker containing 1000 mL of 0.05 mol/L-sodium
115 citrate buffer solution (pH 5.2) with 0.01 g/100 g of sodium azide (final
116 concentration). Each system was heated for 5 min at 70 °C, under stirring, cooled to
117 30 °C and then maintained under constant stirring for 20 h either without or with
118 addition of 0.25 g of hemicellulase. Deionized (Milli-Q™, USA) water was used for
119 all treatments. Insolubles obtained after digestions were separated through filtration
120 under vacuum, with glass fiber filter (Schleicher & Schuell, Dassel, Germany), and
121 cell wall polysaccharides were finally precipitated from each supernatant through
122 ethanol (96 %, v/v) addition (2 volumes). The precipitate was collected through
123 filtration under vacuum using glass fiber filter, washed and, finally, freeze-dried.

124 The fractions obtained are summarized in **Table 1**.

125

126 2.2. Bacterial strains

127 *Lactobacillus plantarum* 8114 (American Type Culture Collection, Rockville,
128 MD, USA), *Bifidobacterium bifidum* ATCC 11863 (MEDICA-TEC, Buenos Aires,
129 Argentina) and *Escherichia coli* 25922 (American Type Culture Collection,
130 Rockville, MD, USA) were used for this study.

131 The specific test strains of *L. plantarum* 8114 and *B. bifidum* 11863 were
132 selected because they were either already established as probiotics or they have
133 potential probiotic properties.

134 All the microorganism cultures were maintained at -80 °C. In the case of
135 *Lactobacillus plantarum* it was used MRS Broth (Difco Laboratories, Sparks, MD,
136 USA) containing 15 % (w/v) glycerol while Tryptic Soy Broth (TSB; Difco
137 Laboratories) containing 15 % (w/v) glycerol was used for *E. coli* and MRS broth
138 (Difco Laboratories, Sparks, MD, USA) supplemented with 0.05 % L-cystein HCl
139 (decrease of oxidation-reduction potential) was used for *Bifidobacterium bifidum*.

140

141 2.3. Prebiotic activity

142 As mentioned before, according to Roberfroid⁴ one of the requirements for the
143 classification of a food ingredient as a prebiotic is the scientific demonstration that it
144 stimulates selectively the growth and/or activity of intestinal bacteria associated with
145 health and wellbeing. This means that the prebiotic activity reflects the ability of a

146 given substrate to support the growth of an organism relative to other organisms and
 147 relative to growth on a non-prebiotic substrate, such as glucose.

148 2.3.1. Prebiotic activity score

149 Huebner et al.¹⁴ established a quantitative score to describe the extent to which
 150 prebiotics support selective growth of lactobacilli and bifidobacteria. This score is
 151 calculated as:

$$\text{Prebiotic activity score} = \left[\frac{(\text{probiotic log CFU/ml on the prebiotic at 48 h} - \text{probiotic log CFU/ml on the prebiotic at 0 h})}{(\text{probiotic log CFU/ml on glucose at 48 h} - \text{probiotic log CFU/ml on glucose at 0 h})} \right] - \left[\frac{(\text{enteric log CFU/ml on the prebiotic at 48 h} - \text{enteric log CFU/ml on the prebiotic at 0 h})}{(\text{enteric log CFU/ml on glucose at 48 h} - \text{enteric log CFU/ml on glucose at 0 h})} \right] \quad \text{Eq (1)}$$

152

153 where CFU means colony forming units.

154 Carbohydrates have a positive prebiotic activity score if they are metabolized
 155 as well as glucose by probiotic strains and are selectively metabolized by probiotics
 156 but not by other intestinal bacteria.

157

158 2.3.2. Prebiotic activity score assay

159 The procedure used is described in **Figure 1**.

160 For prebiotic activity studies, frozen cultures were streaked onto MRS agar for
 161 *L. plantarum* 8114, onto MRS agar supplemented with 0.05 % L-cystein HCl for *B.*
 162 *bifidum* 11863 and onto tryptic soy agar (TSA) for *E. coli* ATCC 25922. Then, *E. coli*
 163 was incubated at 37 °C for 24 - 48 h under aerobic condition, *Lactobacillus plantarum*
 164 and *B. bifidum* were incubated at 37 °C for 24 - 48 h in an anaerobic chamber (Oxoid,
 165 Cambridge, UK) under anaerobic atmosphere (Anaerocult A, Merck, Darmstadt,
 166 Germany). After that, one colony from each plate was transferred into 10 ml of MRS
 167 broth for *L. plantarum* or into 10 ml of MRS broth supplemented with 0.05 % L-
 168 cystein HCl for *B. bifidum* and were incubated overnight in anaerobic conditions. For
 169 *E. coli*, one colony from TSA plate was inoculated into 10 ml of tryptic soy broth
 170 (TSB) and incubated in aerobic conditions for 48 h.

171

172 The assay was performed by adding 1 % (v/v) of an overnight culture of *L.*
 173 *plantarum* to separate tubes containing MRS broth with 1 % (w/v) glucose or 1 %
 174 (w/v) fiber samples. The culture of *B. bifidum* (1 % (v/v)), was added to separate
 tubes containing MRS broth supplement with 0.05% L-cystein HCl and 1 % (w/v)

175 glucose or 1 % (w/v) fiber samples. In both cases, cultures were incubated at 37 °C for
176 48 h under anaerobic atmosphere generation system (Anaerocult A, Merck,
177 Darmstadt, Germany) in an anaerobic chamber (Oxoid, Cambridge, UK). At 0 and 48
178 h of incubation, samples were enumerated in triplicate using the serial dilution method
179 on MRS agar (*L. plantarum*) or MRS agar supplemented with 0.05 % L-cystein HCl
180 (*B. bifidum*) with incubation at 37 °C under anaerobic condition and results were
181 calculated as CFU/mL of culture.

182 *E. coli* culture ATCC 25922 (1 % v/v) was added to separate tubes containing
183 M9 Minimal Medium broth¹⁵ with 1% (w/v) glucose or 1% (w/v) fiber samples and
184 incubated at 37 °C for 48 h in aerobic conditions as described by Huebner et al.^{14,16}
185 and Marotti et al.⁶ At 0 and 48 h of incubation, inoculated samples were enumerated
186 in duplicate on TSA plates with incubation at 37 °C in aerobic conditions. The results
187 were expressed as CFU/mL of culture.

188 Each assay was replicated a minimum of three times.

189

190 2.4. Modelling of the microbial growth

191 Cell counts were evaluated by plating in triplicate after 12, 24, 36, 48 and 60 h
192 of fermentation at 37 °C. Samples (1.0 mL) were added to 9.0 mL of 0.1 g/100 g
193 sterile peptonated water; then, appropriate dilutions were made. Subsequently, *L.*
194 *plantarum* 8114 was plated into MRS Agar and incubated in anaerobic conditions at
195 37 °C. *B. bifidum* 11863 was counted in MRS Agar supplemented with 0.05 % L-
196 cystein HCl with incubation at 37 °C under anaerobic conditions. Incubation was
197 performed for 60 h.

198 *L. plantarum* and *B. bifidum* counts were mathematically modeled for better
199 understanding the behavior of the cultures in the presence of the different fractions of
200 interest. It was used the Gompertz model which is one of the most recommended
201 models^{17,18} and is expressed through the following equation:

202

$$203 \log N = a + c \cdot \exp(-\exp(-b \cdot (t - m))) \quad \text{Eq. 2}$$

204

205 where **log N** is the decimal logarithm of microbial counts (log(CFU/mL)) at time *t*; **a**
206 is the asymptotic log count as time decreases indefinitely which is approximately
207 equivalent to the log of the initial level of bacteria (log (CFU/mL)); **c** is the log count

208 increment or number of log cycles of growth as time increases indefinitely (log
209 (CFU/mL)); **b** is the relative maximum growth rate at time **m** (1/days); **m** is the time
210 required to reach the maximum growth rate (days). Using these parameters, the
211 specific growth rate $\mu = b.c/e$ with $e = 2.7183 (\log(\text{CFU/mLdays}^{-1}))$, lag phase
212 duration (LPD = $m-(1/b)$) (days) and the maximum population density, MPD = $a+c$
213 (log (CFU/mL) can be evaluated.

214

215 2. 5. Statistical analysis

216 Results of experiments are informed as mean \pm standard deviation of three
217 independent determinations. One-way analysis of variance (ANOVA) followed by
218 Duncan's new multiple range tests were used to compare the mean values (α : 0.05).

219 All statistical analyses were performed with SYSTAT INC, version 12.0
220 (Systat Software Inc., San Jose, CA).

221

222 3. Results and Discussion

223 Fissore et al.¹³ informed that the fractions enriched in soluble fiber studied in
224 the present research are constituted by 72.0 - 96.8 g/100g of carbohydrates, 1.8 - 9.2
225 g/100g of proteins and contain phenolic compounds (2.1 - 8.2 g/100g). Carbohydrates
226 comprise uronic acids (14.0 - 18.2 g/100g), neutral sugars (0.8 - 44.3 %) of pectins,
227 and inulin (38.0 - 55.0 %). The highest inulin contents were observed for all fractions
228 isolated in the absence of enzymatic treatment (fractions B, D and F). The lowest
229 degree of methylation of pectin was observed for the fraction isolated from stem in
230 the presence of hemicellulase (fraction A). The lowest protein and phenol contents
231 were observed for fractions isolated from bracts (fractions C and D) (**Table 2**).

232

233 3. 1. Kinetic behavior of the *Lactobacillus plantarum* 8114 and *Bifidobacterium* 234 *bifidum* 11863 growth in different fibers

235 When studying the substrate requirements and specificities of individual
236 Bifidobacterial and Lactobacillus strains, two factors are especially important. The
237 first is the rate at which an organism can grow on a particular carbon source, as this
238 will influence its ability to compete with other bacteria in the colon.¹⁹ The other is the
239 extent to which the substrate is converted into bacterial mass, because cell numbers
240 will affect the degree of pre- or probiotic activity. For this reason, it is important the

241 study of the kinetic behavior of the probiotic bacteria *Lactobacillus plantarum* 8114
242 and *Bifidobacterium bifidum* 11863 in the different substrates.

243 **Figure 2** shows *Lactobacillus plantarum* 8114 (Panel a) and *Bifidobacterium*
244 *bifidum* 11863 (Panel b) growth on the different fractions of dietary fiber during
245 incubation at 37 °C for a maximum period of 60 h. Full lines represent the
246 mathematical modeling of data to the Gompertz equation. As can be observed, a good
247 agreement was achieved between the model and the experimental data; the parameters
248 obtained are shown in **Table 3**.

249 In the case of *Lactobacillus plantarum* 8114 strains, the highest specific
250 growth rate (μ : 0.16 1/h) was observed for fraction A, indicating that a high rate of
251 cell proliferation occurred on this carbon source within a short period of incubation
252 (**Table 3**). For B, C, D E and F fractions, the specific growth rate (μ) was similar to
253 the one observed for glucose (0.09 1/h). The maximum population density (MPD) was
254 similar for glucose and fraction A and these were the higher values observed (9.88 -
255 10.11 log CFU/mL) while for other fractions the MPD values were in the range 8.47 -
256 9.18 log CFU/mL. The lag phase duration (LPD) for the fractions ranged from 11.62
257 to 21.62 h and for glucose it took a significantly lower value of 4.90 h.

258 In the case of *Bifidobacterium bifidum* 11863, the highest specific growth rate
259 was obtained for fractions A and B (0.08 - 0.09 1/h) and this rate doubled the value
260 observed for MRS broth with glucose (0.04 1/h) but differences were not significant
261 for the growth on fraction A and glucose. The other fibers showed specific growth
262 rates of 0.05 - 0.07 1/h and differences between fibers were not statistically significant
263 ($p > 0.05$). MPD values ranged between 8.32 - 9.07 log (CFU/mL) for different
264 fractions while for glucose, the MPD value was 8.65 log (CFU/mL). The lag phase
265 duration (LPD) showed significant variation for the different fractions ranging
266 between 8.83 and 21.16 h and a value of 4.62 h was observed for glucose (**Table 3**).

267 Values obtained for specific growth rate are similar to those informed by
268 Marotti et al.⁶ for *Lactobacillus* and *Bifidobacterium* on soluble fibers from modern,
269 old durum and ancient type wheat varieties. Hernandez-Mendoza et al.²⁰ reported
270 higher specific growth rates and similar MPD for *Lactobacillus reuteri* and
271 *Bifidobacterium bifidum* inoculated into a reconstituted whey containing sucrose and
272 pectin in order to prepare a fermented probiotic product.

273 It can be concluded that *L. plantarum* 8114 showed a higher specific growth
274 rate on fraction A than on glucose. Specific growth rate values were higher for this
275 strain than for *Bifidobacterium bifidum* 11863 although differences were not
276 statistically significant ($p>0.05$).

277

278 3. 2. Growth of *Lactobacillus plantarum* 1814, *Bifidobacterium bifidum* 11863 and 279 *Escherichia coli* 25922 on fractions enriched in soluble fiber

280 One of the characteristic properties of a prebiotic substrate is that it should
281 stimulate selectively the growth and/or activity of intestinal bacteria associated with
282 health and wellbeing. Thus, it was studied the increase in population cell number for
283 strains of *Lactobacillus* and *Bifidobacterium* following 48 h growth on 1 % (w/v)
284 glucose or on 1 % (w/v) fraction enriched in soluble fiber and the same procedure was
285 used to study the growth of *E. coli* 25922 which was chosen to represent the enteric
286 portion of the commensal flora. The results are shown in **Table 4**.

287 For *Lactobacillus* strain, increase in cell density (CFU/mL) on fractions B, C,
288 D, E and F was significantly lower (1.57-2.11) than cell density increase on glucose
289 (3.00). The increase in cell density of *L. plantarum* 8114 for fiber A (2.94) and for
290 glucose were similar. In the case of *B. bifidum* 11863, a significantly higher ($p<0.05$)
291 increase in cell density was observed when fibers A, B, C, D or F were present (1.90-
292 2.03) than when glucose (1.60) was in the media. Growth of *E. coli* 25922 on all the
293 fractions studied was significantly lower (0.56-0.62) than the growth on glucose
294 (1.47) as can be observed in **Table 4**.

295

296 3.3. Prebiotic activity score

297 Prebiotic activity scores for *Lactobacillus plantarum* 8114 and
298 *Bifidobacterium bifidum* 11863 shown in **Table 5** were derived from the cell density
299 values of **Table 4** through the use of **Eq. 1**. All scores calculated were positive. The
300 higher the score, the higher the relative growth of the probiotic and/or the lower the
301 relative growth of the *E. coli*, which indicates a higher and more selective use of
302 prebiotic in relation to glucose by the probiotic microorganism and/or a limited use of
303 prebiotic in relation to glucose by *E. coli*.

304 The highest prebiotic activity score was observed for *Bifidobacterium bifidum*
305 grown in MRS broth and with fiber B added (0.87) and the score for the other fibers
306 were not significantly different ($p>0.05$).

307 For *Lactobacillus plantarum*, the highest prebiotic score was observed for the
308 microorganism grown on MRS broth with fiber A added (0.58). Lower scores were
309 observed when *L. plantarum* was grown in the presence of fibers C, F, D, E and B
310 (0.31, 0.24, 0.19, 0.16 and 0.14, respectively) although differences were not
311 statistically significant ($p>0.05$).

312 As can be observed in **Table 5**, there are significant differences ($p<0.05$) in
313 prebiotic activity scores between the two strains grown on fractions B, C, D, E and F,
314 being the values for *Lactobacillus plantarum* lower than those for *Bifidobacterium*
315 *bifidum*. This indicates that differences in their metabolic capacity apparently existed.
316 The utilization of different fractions by the studied bacteria requires the presence of
317 specific hydrolysis and transport systems and its presence or absence may be the
318 cause for the different prebiotic scores observed.¹⁴

319 The capacity of Lactobacilli and Bifidobacteria to utilize a diverse range of
320 dietary carbohydrates has been previously informed and the literature link this
321 capacity to a metabolic adaptation to a complex carbohydrate-rich gastrointestinal
322 tract environment. According to Pokusaeva et al.²¹, for an average individual the
323 human gastrointestinal tract (GIT) is a natural habitat for approximately 10^{11} – 10^{12}
324 microorganisms per gram of luminal content, collectively forming the gut microbiota
325 with a total biomass of more than 1 kg in weight. The total number of bacterial
326 species that may be contained within the intestinal microbiota, ranges from
327 approximately 500 to 1,000 distinct bacterial species to between 15,000 and 36,000
328 different species. Lactobacilli and Bifidobacteria are among the prevalent groups
329 thought to exert health-promoting actions in the GIT. Bifidobacteria can utilize a
330 diverse range of dietary carbohydrates that escape degradation in the upper parts of
331 the intestine, many of which are plant derived oligo- and polysaccharides. Different
332 bifidobacterial strains may possess different carbohydrate utilizing abilities. The gene
333 content of a bifidobacterial genome reflects this apparent metabolic adaptation to a
334 complex carbohydrate-rich gastrointestinal tract environment as it encodes a large
335 number of carbohydrate-modifying enzymes and this is a subject of actual study. The
336 capacity of individual strains and species of Lactobacilli for carbohydrate metabolism

337 differs substantially. This metabolic diversity conforms to the phylogenetic diversity
338 in the genus *Lactobacillus*. Several species like *L. acidophilus*, *L. casei*, and *L.*
339 *plantarum* metabolize a large diversity of different carbon sources, including all major
340 categories of oligo- and polysaccharides. Oligosaccharides are preferentially
341 metabolized by phosphotransferase/phospho-glycosyl hydrolase systems and
342 oligosaccharide metabolism is repressed by glucose. Other species exhibit more
343 restricted carbohydrate fermentation patterns being an extreme the “nothing but
344 maltose or sucrose” diet of several strains of *L. sanfranciscensis*. In this group of
345 strains, oligosaccharides are preferentially metabolized by permease/phosphorylase
346 systems and oligosaccharide metabolic enzymes are not repressed by glucose²². Both
347 groups are represented in intestinal habitats (e.g., *L. acidophilus* and *L. reuteri*) as
348 well as food fermentations (e.g., *L. plantarum* and *L. sanfranciscensis*) and actual
349 studies of carbohydrate consumption in model substrates, and in food or intestinal
350 ecosystems are trying to improve the understanding on these phenomena.²²

351 Parkar et al.²³ reported gut health benefits exerted by kiwifruit pectins.
352 Dongowski et al.²⁴ investigated the degradation, metabolism, fate, and selected effects
353 of pectin in the intestinal tract of rats. They observed that total anaerobic and
354 Bacteroides counts were greater in groups fed with pectin and that they presented a
355 higher concentration of short chain fatty acids (SCFA) in cecum and feces. During *in*
356 *vitro* fermentation of pectin with fecal flora from rats, unsaturated oligogalacturonic
357 acids appeared as intermediate products. With increasing degree of methylation, the
358 formation rate of SCFA decreased in the cecum of conventional rats. Low methoxyl
359 pectins was fermented faster than high methoxyl pectins *in vivo* and *in vitro*.

360 It has been reported that both inulin and oligofructose are effective prebiotics
361 due to the stimulation of colonic bifidobacteria. Because of their recognized prebiotic
362 properties, both are increasingly used in new food product developments such as
363 drinks, yoghurts, biscuits. Bifidobacteria can inhibit gut pathogen growth producing
364 the fortification of the gut flora to resist acute infections.^{25,26,27,28}

365 It can be concluded that fraction A presented the best performance concerning
366 the growth of both strains. According to previous cited bibliography, it might be the
367 content of inulin and of pectin of low degree of methylation, the compositional
368 reasons for its selective stimulation of *Lactobacillus plantarum* 8114 and
369 *Bifidobacterium bifidum* 11863 growth.

370

371 **4. Conclusions**

372 Dietary fiber fractions studied showed, in general, a potential capacity for
373 selectively stimulate the growth of intestinal bacteria associated with health. Fraction
374 isolated from artichoke stem with the use of a heat pre-treatment and hemicellulase
375 followed by ethanol precipitation (fraction A) had the highest prebiotic activity score
376 for both strains since it was determined:

- 377 - the highest specific growth rate of *Lactobacillus plantarum* 8114 on this fraction
- 378 with respect to glucose,
- 379 - a similar population density achieved by *Lactobacillus plantarum* 8114 and
- 380 *Bifidobacterium bifidum* 11863 when grown on this fraction and on glucose,
- 381 - the smaller increase in cell density observed for *Escherichia coli* 25922 on this
- 382 fraction with respect to glucose,
- 383 - the smaller increase in cell density observed for *Escherichia coli* 25922 in
- 384 comparison to that of *Lactobacillus plantarum* 8114 and *Bifidobacterium bifidum*
- 385 11863 when grown on this fraction.

386 This behavior might be attributed to the inulin and low methoxyl pectin contents of
387 fraction A.

388 Other fractions also produced high prebiotic activity scores for
389 *Bifidobacterium bifidum* 11863 but they showed lower prebiotic activity scores for
390 *Lactobacillus plantarum* 8114.

391 The potential of fraction A to promote the growth of both tested strains in the
392 gastrointestinal tract is promising. It is necessary to perform additional studies in
393 order to evaluate the resistance of these fractions to different pHs and enzymes
394 present in the human gastrointestinal tract and to analyze their gastrointestinal
395 absorption and fermentation by the intestinal microflora where the competition for
396 nutrients may influence bacterial survival, colonization and metabolic activity in the
397 host.

398

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403 11220090100531 and 11220120100507).

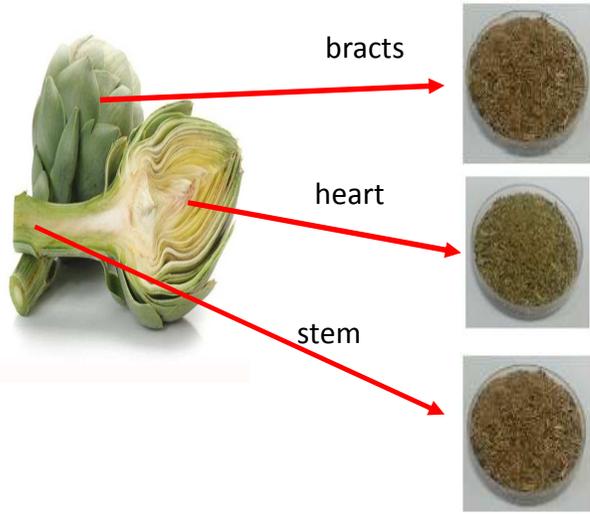
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405 **References**

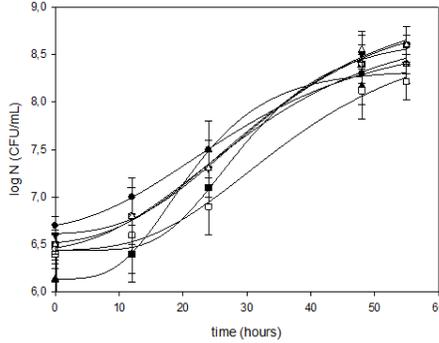
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Cell Wall Material



Fractions enriched in soluble fiber



growth of intestinal bacteria associated with health

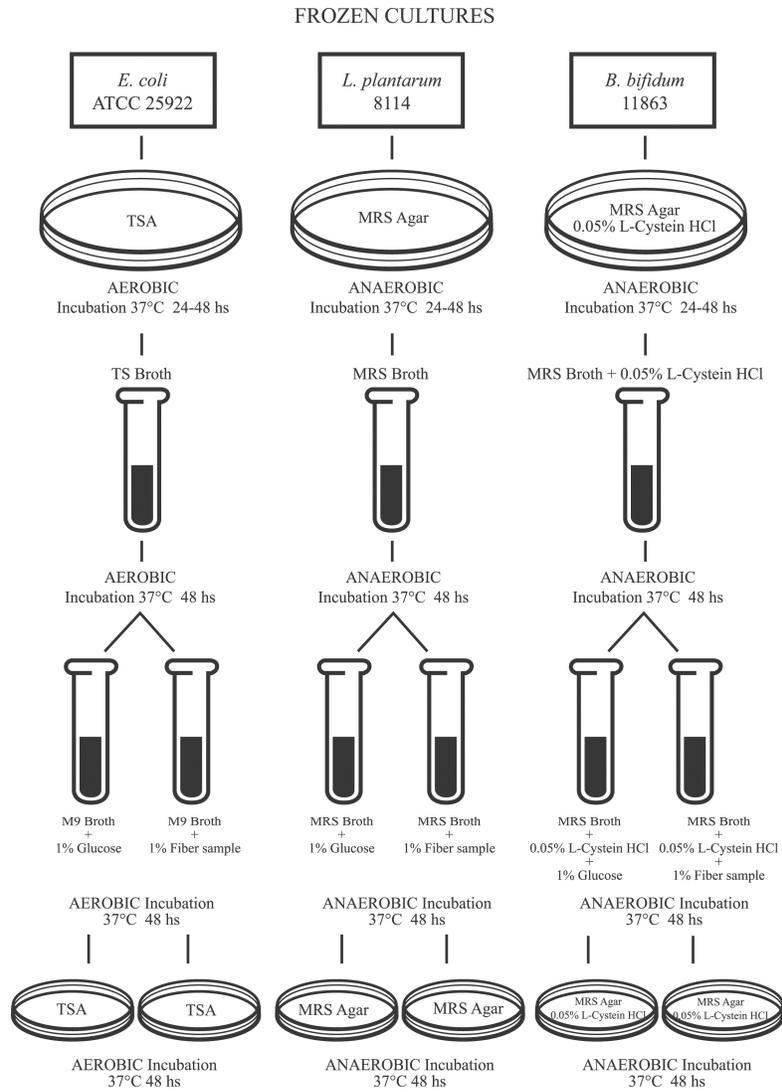


Figure 1: Flow-graph of method used for prebiotic activity score assay

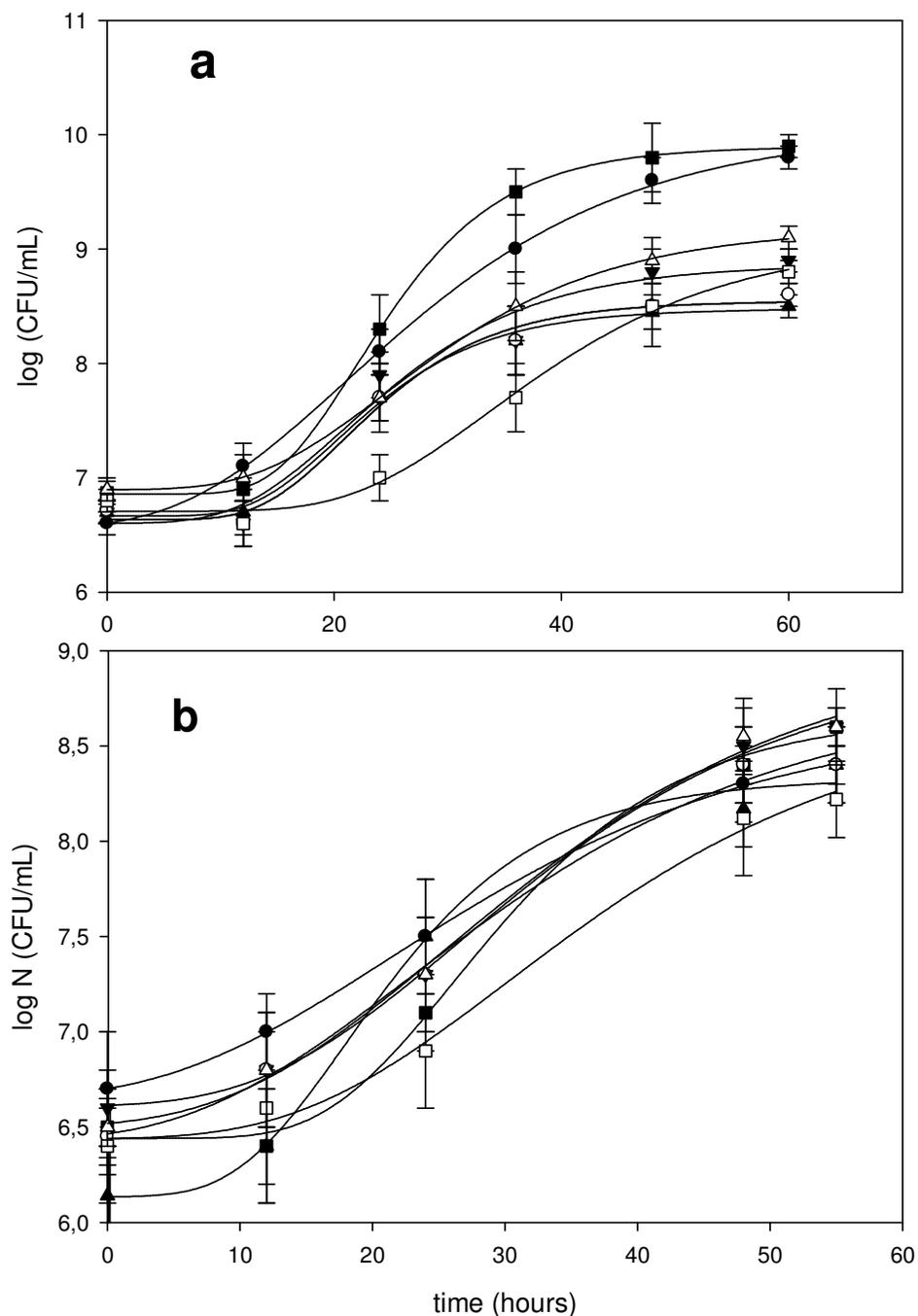


Figure 2. Application of Gompertz model to experimental data of *Lactobacillus plantarum* 8114 (a) and *Bifidumbacterium bifidum* (b) growth in different type of fibers: ● MRS broth with glucose (1% w/v), ■ MRS broth with fiber A (1% w/v), ▲ MRS broth with Fiber B (1% w/v), ▼ MRS broth with Fiber C (1% w/v), ○ MRS broth with Fiber D (1% w/v), □ MRS broth with Fiber E (1% w/v) and △ MRS broth with Fiber F (1% w/v).

Table 1. Different fractions obtained from the treatment of artichoke cell wall material (CWM)

Fraction	CWM from artichoke	Treatment with hemicellulase
A	stem	+
B	stem	-
C	bracts	+
D	bracts	-
E	heart	+
F	heart	-

Table 2. Chemical composition of the fractions enriched in soluble fibers and isolated from bracts, hearts and stems of artichoke¹

Fraction ²	Total carbohydrates (g per 100 g of fraction)	Uronic acids (g per 100 g of fraction)	Inulin (g per 100 g of fraction)	Neutral sugars (g per 100 g of fraction)	Protein (g per 100 g of fraction)	Total phenolics (g per 100 g of fraction)	DM ³
A	76.0± 7.0	15.0 ± 0.1	46.0 ± 0.4	15.0	6.8 ± 0.1	6.9 ± 0.2	15
B	72.0 ± 6.0	18.2 ± 0.2	53.0 ± 0.1	0.8	9.2 ± 0.2	4.8 ± 0.2	33
C	83.3 ± 0.1	14.0 ± 0.1	38.0 ± 1.0	31.3	1.8 ± 0.5	3.1 ± 0.2	31
D	76.0 ± 7.0	14.2 ± 0.2	44.7 ± 0.3	17.1	2.7 ± 0.3	2.1 ± 0.1	31
E	96.8 ± 0.3	14 ± 1	38.5 ± 0.2	44.3	7.9 ± 0.1	8.2 ± 0.2	39
F	79.0 ± 6.0	15.1 ± 0.1	55.0 ± 0.1	8.9	5.8 ± 0.1	4.0 ± 0.1	58

¹ Fissore et al (2014).

² A: fraction obtained form artichoke stem CWM with hemicellulase. B: fraction obtained form artichoke stem CWM with no enzyme addition. C: fraction obtained from artichoke bracts CWM with hemicellulase. D: fraction obtained from artichoke bracts CWM with no enzyme addition. E: fraction obtained from artichoke heart CWM with hemicellulase. F: fraction obtained from artichoke heart CWM with no enzyme addition. CWM: cell wall material.

³DM: Degree of methylation. Ratio between moles of methanol and moles of GalA (uronic acids) per 100 g of sample.

Table 3. Gompertz parameters: specific growth rate (μ), maximum population density (MPD) and lag phase duration (LPD) for *Lactobacillus plantarum* 8114 and *Bifidumbacterium bifidum* 11863 growth in MRS broth with glucose or different fractions isolated from artichoke

<i>Lactobacillus plantarum</i> 8114			
Substrate	μ (1/h)	LPD (h)	MPD Log (CFU/mL)
Glucose (MRS)	0.09±0.009A	4.90±0.98A	10.11±0.26A
Fraction A	0.16±0.06B	14.75±0.36BD	9.88±0.05B
Fraction B	0.09±0.02A	13.39±1.88B	8.47 ± 0.18C
Fraction C	0.09±0.04A	11.62±6.07B	8.86 ±1.67BC
Fraction D	0.09±0.03A	13.53±3.42B	8.54±0.32C
Fraction E	0.07±0.02A	21.62±4.39CD	9.06±0.40C
Fraction F	0.08±0.01A	13.27±0.45B	9.18±0.05C
<i>Bifidumbacterium bifidum</i> 11863			
Glucose (MRS)	0.04±0.001A	4.62±1.23A	8.65±0.10A
Fraction A	0.08±0.05A	16.10±4.25BD	8.66±0.45A
Fraction B	0.09±0.03B	9.70±2.80B	8.32 ±0.33A
Fraction C	0.05±0.03A	13.54±3.01B	8.83 ±0.90A
Fraction D	0.05±0.02A	10.20±4.71B	8.63±0.44A
Fraction E	0.07±0.02A	21.16±2.71CD	9.07±0.40A
Fraction F	0.05±0.03A	8.83±5.20B	9.02±0.78A

Capital letters are used to describe differences in parameters in each column. Different letters correspond to significant differences between values.

Table 4. Increase in cell density between time 0 and time 48 h, reported as log(CFU/ mL) standard deviation, for bacterial cultures grown on glucose or on different fractions isolated from artichoke

Substrate	<i>L. plantarum</i> 8114	<i>Bifidobacterium</i> <i>bifidum</i> 11863	<i>E. coli</i> 25922
Glucose	3.00±0.15A a	1.60 ±0.10A b	1.47±0.06A b
Fraction A	2.94±0.19A a	1.90±0.09B b	0.58±0.11B c
Fraction B	1.57±0.20B a	2.03±0.09B b	0.56±0.10B c
Fraction C	2.11±0.13B a	1.90±0.11B a	0.57±0.10B b
Fraction D	1.84±0.21B a	1.95±0.09B a	0.60±0.08B b
Fraction E	1.70±0.23B a	1.70±0.08A a	0.59±0.09B b
Fraction F	2.00±0.11B a	2.00±0.11B a	0.62±0.09B b

Capital letters are used to describe differences in cell density in each column.

Lowercase letters are used to describe differences in cell density in each row.

Different letters correspond to significant differences between values.