



Bioaccessibility and bioavailability of phenolic compounds in bread: a review

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Thank you for your effort in reviewing this submission. It is only through the continued service of referees that we can maintain both the high quality of the publication and the rapid response times to authors. We would greatly appreciate if you could review this paper in **14 days**. Please let us know if that will not be possible.

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Best wishes,

Philippa Hughes
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DEGLI ALIMENTI E DEL FARMACO

To Prof. Laura Bravo-Clemente,
Institute of Food Science, Technology and Nutrition,
Madrid, Spain

Parma, April 12th 2017

Dear Editor,

following the invitation by the Editorial Office of Food & Function, we are submitting a review manuscript entitled "Bioaccessibility and bioavailability of phenolic compounds in bread: a review" by Angelino and colleagues.

The aim of this review is to provide an overview of the literature related to the bioaccessibility and bioavailability of phenolic compounds in bread. We tried to focus mainly on the potential strategies to improve phenolic bioaccessibility and bioavailability and to the main findings of in vitro and in vivo studies investigating these strategies applied to breads.

We confirm that the paper is not under submission to other journals and that all the authors read and approved the final manuscript.

We hope that you will consider our work for publication in Food & Function.

Regards,



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DIPARTIMENTO DI SCIENZE
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1 **Bioaccessibility and bioavailability of phenolic compounds in bread: a review**

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29 **ABSTRACT**

30 Cereal-based products, like breads, are a vehicle for bioactive compounds, including polyphenols. The health
31 effects of polyphenols like phenolic acids (PAs) are dependent on their bioaccessibility and bioavailability.
32 The present review summarizes the current understanding of potential strategies to improve phenolic
33 bioaccessibility and bioavailability and the main findings of *in vitro* and *in vivo* studies investigating these
34 strategies applied to breads, including the use of raw ingredients with greater phenolic content and different
35 pre-processing technologies, such as fermentation and enzymatic treatment of ingredients. There is
36 considerable variability between *in vitro* studies mainly resulting from the use of different methodologies,
37 highlighting the need for standardization. Of the few *in vivo* bioavailability studies identified, acute, single-
38 dose studies demonstrate that modifications to selected raw materials and bioprocessing of bran could
39 increase the bioavailability, but not necessarily net content, of bread phenolics. The two medium term
40 identified dietary interventions also demonstrated greater phenolic content resulting from modification of
41 raw materials used. Overall, findings suggest that several strategies can be used to develop new bread
42 products with greater phenolic bioaccessibility and bioavailability. However, due to the large variability and
43 the few studies available, further investigations are required to better determine the usefulness of these
44 innovative processes.

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46 **KEYWORDS:** bread, bioaccessibility, bioavailability, phenolic compounds.

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58 **INTRODUCTION**

59 Cereal-based products are the most common staple foods globally. Among the wide range of products, bread
60 is one of the most consumed. The estimated bread consumption has been reported to be over 100 g per day
61 (equivalent to approximately 3 slices per day) in many countries¹⁻³, therefore bread is an important
62 contributor to daily energy intake⁴.

63 Bread products differ widely in shape, size, texture, and sensory characteristics. Part of these differences are
64 ascribable to the type of cereal used for bread-making, which can include rye, barley, oat, and wheat, the
65 latter of which is the most commonly used due to its gluten content, which contributes to good sensory
66 characteristics. Differences can also result from the addition of ingredients, such as seeds, olives and nuts, as
67 well as differences in the bread-making process, such as temperature and the use of yeast versus sourdough.
68 Regardless of these differences, bread is generally characterized by a high carbohydrate and protein content,
69 but it is also a rich source of vitamins (mainly from the B-vitamin group) and minerals (such as iron,
70 calcium, phosphorus, zinc, potassium, and magnesium).

71 Many bread products are also a good source of bioactive compounds, including fibre and other
72 phytochemicals, specifically those made with wholegrains that consist of the intact, ground, cracked or
73 flaked kernel after the removal of inedible parts such as the hull and husk. Only in wholegrain products are
74 the principal anatomical components, including the starchy endosperm, germ and bran, present in the same
75 relative proportions as in the intact kernel⁵. In the outer layers of the kernels, where the bran is found, there is
76 a high content of bioactive compounds⁶.

77 The consumption of whole grains has been associated with the prevention of chronic diseases, including
78 cardiovascular disease and diabetes^{7,8}. Therefore, clinical practice guidelines and dietary guidelines
79 recommend choosing wholegrain products⁹⁻¹², which are rich in bioactive compounds, over refined products,
80 in which bioactive compounds are present only in small amounts due to the removal of the seed external
81 layers during milling.

82 Wholegrain bread products are rich in fibre, particularly insoluble fibre, for which bran represents one of the
83 main sources. Fibre from bread products mainly includes arabinoxylans, a hemicellulose found in plant cell
84 walls and that represent the major component of dietary fibre in cereal grains. The wheat grain also contains

85 aleurone as a monolayer of cells overlying the endosperm, which is rich in fibre and phenolic compounds.
86 Furthermore, breads can be also rich in soluble fibres, like those made with oat and barley as good sources of
87 β -glucans, which are well known to reduce post-prandial blood glucose and blood cholesterol¹³⁻¹⁵, risk
88 factors in the development of coronary heart disease (CHD)¹⁶.
89 Similarly to other cereal-based products, wholemeal bread is generally a good source of phenolic
90 compounds, mainly as esters bound to arabinoxylans¹⁷, with a minor contribution of soluble free or
91 conjugated compounds¹⁸. Polyphenols exist as secondary metabolites in several different plants, in which
92 they can act as a defence mechanism against parasites and toxic compounds^{19,20}. Phenolic compounds are
93 widely diffused in all plant foods including fruits, vegetables and beverages (tea and coffee), the
94 consumption of which may lead to a phenolic intake of ~1000 mg per day, in a typical American diet²¹.
95 Bread products contribute to this daily phenolic intake, especially when they include bran.
96 Cereal grains constitute a good source of phenolic acids (PAs), in addition to alkylresorcinols and lignans.
97 PAs can be divided in two groups, hydroxycinnamic and hydroxybenzoic acids, deriving from the
98 hydroxylation of the cinnamic or benzoic acid moiety. Hydroxycinnamic acids are the most abundant PAs
99 and chiefly consist of ferulic acid (FA), *p*-coumaric acid (CA), caffeic acid, and sinapic acid (SA).
100 Hydroxybenzoic acid derivatives include *p*-hydroxybenzoic, protocatechuic, vanillic, syringic and gallic
101 acids.
102 Polyphenols are not included in the category of micronutrients, as they are not essential for the maintenance
103 of vital functions. However, several studies indicate that phenolic compounds might be responsible for part
104 of the beneficial effects associated with the consumption of plant-based foods, such as the association
105 between fruit and vegetable intake and reduced CVD risk²². In particular, *in vitro* studies have demonstrated
106 the involvement of polyphenols and their metabolites in several features linked to prevention of
107 inflammation, oxidative stress and many other recognised pathophysiological processes²³⁻²⁵. Furthermore,
108 over the past 20 years, epidemiological studies have demonstrated that the consumption of polyphenol-rich
109 foods, such as fruits, vegetables, cereals, coffee and cocoa, is inversely associated with the risk of many
110 chronic diseases. The first epidemiological study focusing on the protective role of polyphenols on CHD
111 found a 42% reduction in relative risk of CHD mortality when comparing the highest tertile of flavonoid
112 intake to the lowest²⁶. Several other epidemiological studies followed, including the Iowa Women's Health

113 Study (n= 41,836), in which polyphenol intake was inversely associated with inflammation²⁷, and
114 specifically, whole grain polyphenol intake was inversely associated with the incidence of colorectal
115 cancer²⁸.

116 Evidence from human intervention trials on the protective effects of phenol-rich foods against many chronic
117 diseases has been inconsistent, possibly because of differences in food composition, as well as differences in
118 the absorption and metabolism of various phenolic compounds. One of the main issues contributing to the
119 inconsistency in results concerns studies attributing the effects to a single compound or a class of foods,
120 because a single compound can be present in several different foods, and a class of foods can contain
121 mixtures of polyphenols. In addition, the *in vivo* effects of polyphenols are strongly influenced by their
122 bioavailability.

123 Generally, bioavailability is the fraction of an ingested nutrient or compound that reaches the systemic
124 circulation and may be utilized. Thus, it is multifactorial in that it includes gastrointestinal digestion,
125 absorption, metabolism, tissue distribution, and bioactivity of the nutrient/compound. However, due to the
126 difficulty in investigating the bioactivity, bioavailability is commonly considered both the fraction of a
127 compound as well as the metabolite(s) of that compound that reach the systemic circulation²⁹.

128 Bioavailability can be affected by a wide range of factors, not only related to the food (e.g. chemical form of
129 the compound, characteristics of the food matrix), but also to the individual (e.g. gastric emptying, intestinal
130 transit time), resulting in high inter-individual variability^{24,30}.

131 Beginning with ingestion and digestion of a food, the food matrix can influence the bioaccessibility of the
132 phenols because the amount that is released from within the matrix will influence the fraction that is made
133 available for intestinal absorption. Effects on bioaccessibility can be evaluated *in vitro* by simulation of
134 gastric and small intestinal digestion³¹. *In vitro* methods are quick and inexpensive ways to estimate the
135 bioaccessibility of a bioactive compound, including changes resulting from variations in the food matrix and
136 food processing. However, these methods cannot completely measure of the bioavailability of bioactives, as
137 this requires *in vivo* methodologies.

138 The present review summarizes potential strategies, including innovative technologies, that can be applied
139 during the bread-making process in an effort to increase the fraction of phenolic compounds reaching the

140 systemic circulation, and what is currently known about the usefulness of these strategies as assessed in *in*
141 *vitro* bioaccessibility and *in vivo* bioavailability studies.

142

143 **POTENTIAL STRATEGIES TO INCREASE PHENOLIC CONTENT IN BREAD PRODUCTS:** 144 **EFFECTS OF THE BREAD-MAKING PROCESS**

145 Various processing techniques are applied to grains in order to transform the raw materials into finished
146 products with good sensory characteristics and nutritional quality. Since technological processes affect the
147 chemical constituents and physical properties of foods, it is expected they also influence the phenolics within
148 grain products, thus impacting the potential beneficial health effects. The effect of various food-processing
149 methods on phenolic compounds has therefore become an important area of research.

150 A review of the literature has highlighted three main strategies that can be applied to design phenolic-
151 enriched breads: the first approach focuses on the use of raw materials naturally rich in phenolic compounds;
152 the second focuses on the application of bio-processing techniques on raw materials; and the third focuses on
153 the processing conditions that can be applied during bread-making (Table 1).

154 **Raw materials**

155 Whole grains are a good source of phenolic compounds, mostly concentrated in the bran, but levels of
156 phenolics in the final products can vary widely depending on the raw materials and on the pre-processing
157 techniques. In addition to whole wheat, barley, and rye, minor cereals (e.g. sorghum, millets;³²), pigmented
158 grains³³⁻³⁵, and ancient grains (e.g. eikorn, emmer³⁶ and pseudocereals like buckwheat, quinoa and
159 amaranth)³⁷ represent a good source of phenolics, thus their use in bread products has increased in the
160 marketplace. Since phenolic compounds are present in the external layers of the kernel, adding bran fractions
161 to refined flour is one of the most common trends to enhance phenolic content in bread products.

162 **Pre-processing techniques**

163 Besides using wheat bran and whole-grain flour, several modifications to pre-processing techniques can be
164 used to influence phenolic content in bread products. A variety of fractionation methods, including both wet
165 extraction and dry fractionation, have been developed for producing milling fractions that are concentrated in
166 phenolic compounds. Among fractionation methods, debranning (also named pearling) is the most widely
167 used. It has been traditionally used as a tool to enhance both hygienic and technological performances of

168 milled flours^{38,39}. More recently, debranning has been demonstrated as an effective strategy to produce bran
169 fractions rich in aleurone particles⁴⁰, which are particularly rich in phenolics, thus recovering the bioactive
170 compounds that are concentrated in the external layers of grain kernels⁴¹⁻⁴³.

171 Regarding to physical treatments, air classification technology is an effective way to separate grain flours
172 into fractions with different sizes, properties, and chemical composition, such as protein, starch, and dietary
173 fibre. When applied to phenolic-rich material, it is a good technique to select fractions with a high content of
174 phenolic compounds⁴⁴.

175 Micronization, also known as ultrafine grinding, is a mechanical treatment, used to change or damage the
176 fibre matrix, causing some phenolics which were linked or embedded into the matrix to be exposed so that
177 the total phenolic content in bran increases⁴⁵, likely due to an increase in extractability.

178 Lastly, biotechnological processes (*i.e.* germination, fermentation, and enzymatic treatments) have been used
179 to improve the PA content in bran. Germination is the process by which a plant grows from a seed. During
180 germination, high levels of hydrolytic enzymes, such as amylases and proteases, accumulate in the cereal
181 seed, so that the insoluble endosperm starch and protein reserves are hydrolyzed into soluble forms that can
182 be transported to the embryo to meet the needs of the growing plant. A recent review on the effects of grain
183 germination concluded that during this process a net increase in total phenolic content and total antioxidant
184 capacity⁴⁶ is observed. It is also thought that germination may increase the extractability of polyphenolic
185 compounds, by releasing bound polyphenols, therefore making them more soluble in extraction solvents.

186 Fermentation is another beneficial pre-processing technique which effectively releases phenolics from the
187 bran of various grains^{47,48}. The enzymes produced by the added microorganisms have the potential to release
188 insoluble bound PAs from bran and thereby improve their bioaccessibility and potential bioavailability⁴⁹. In
189 the case of sourdough fermentation, the effect of the reduction in pH is also important^{50,51}. The lower pH
190 during sourdough fermentation favors the activity of hydrolases and can contribute to chemical disintegration
191 of arabinoxylans, and to extensive hydrolysis of both esters and glycosides of PAs^{50,51}.

192 Combining fermentation with germination results in an additive effect, since germination results in a higher
193 amount of fermentable sources (sugars and nitrogen) and both increase the concentration of cell wall
194 degrading enzymes, all contributing to increased bioaccessibility of PAs⁴⁷.

195 A third biotechnological process which can be applied during pre-processing is enzymatic treatment,
196 whereby grains or bran are pre-treated with enzymes in a liquid environment. Enzymatic treatment has been
197 reported to free PAs from fibre esters⁵², improving the bioavailability of these compounds⁵³. Enzymes (e.g.
198 xylanases) are also commonly used in the baking industry, as part of dough conditioners, to improve dough
199 property, baking quality, and shelf-life⁵⁴.

200 **Bread-making process**

201 In addition to the formulation and pre-processing of bread products, the bread-making process also
202 influences the content and bioavailability of phenolic compounds in the final bread product.

203 Bread-making includes several fundamental operations, namely mixing and kneading, fermentation or
204 leavening, and baking, which are indispensable for producing an attractive end product. During mixing,
205 ingredients are evenly distributed and blended. In wheat breads, interaction with water leads to significant
206 structural changes in proteins, resulting in gluten formation; a three-dimensional network structure resulting
207 in a cohesive, completely homogenous, non-sticky mass with well-defined rheological characteristics. These
208 attractive properties depend on the procedure applied and equipment used, as well as on the presence of
209 components, such as phenolics, that may negatively affect gluten viscoelasticity. For example, phenolic
210 compounds can form complexes with proteins, via hydrogen bonding between the hydroxyl groups of the
211 phenols and the carbonyl group of the peptide residue⁵⁵⁻⁵⁷.

212 Studies have demonstrated that dough mixing causes an overall decrease in total PAs, such as bound FA, SA
213 and CA, in various grains^{50,58,59} reaching up to 50%. However, free FA has been demonstrated to increase
214 significantly in one study showing up to five times the initial level, suggesting that mixing may also facilitate
215 the release of bound phenolic compounds into free and more bioaccessible forms^{35,58}.

216 Various mechanisms have been proposed to explain the overall decrease in PAs resulting from dough
217 mixing. High-speed mixing breaks protein disulfide bonds and creates thiol free radicals in gluten, which
218 then react with reducing compounds, like PAs, in flour⁶⁰. Considering the proposed effect of mixing on the
219 formation of bonds between phenolics and proteins, a decrease in phenolic content in various reports may be
220 more accurately described as a reduction in their bioaccessibility and thus extractability⁶¹.

221 Another proposed effect of mixing on phenolics is the hydrolysis of oxidative enzymes such as oxygenase
222 and peroxidase, that are present in flours, which become active when water is added and thus decrease the

223 amount of phenolics like FA⁶².

224 The leavening and fermentation process increases the original volume of the bread and creates a porous
225 structure, through the action of a leavening agent, usually baker's yeast (i.e. *Saccharomyces cerevisiae*),
226 which converts the fermentable sugars present in the dough into ethanol and CO₂.

227 The fermentation and leavening process may contribute to an increase in PA bioavailability. Two
228 mechanisms for the fermentation-induced increase in bioaccessibility and bioavailability of phenolic
229 compounds during bread-making have been proposed: i) via the structural breakdown of the cell wall matrix
230 by degrading enzymes present in both grains and microbes activated by the leavening agent⁶³; and ii) via the
231 synthesis or enzymatic transformation of various bioactive compounds⁴⁷. However, studies investigating PA
232 content in fermented dough are not consistent. This inconsistency is likely due to differences in the enzymes
233 produced from yeast or other microorganisms and native enzymes present in various types of grains. As an
234 example, rye has been described to have much more native enzymatic activity compared to wheat^{35,50,51,59}. In
235 addition, fermentation conditions, particularly temperature, pH, and duration, are contributing factors to PA
236 content. With regards to the fermentation time, prolonged fermentation increases the number of bonds
237 broken between PAs and dietary fibre, thus increasing the bioaccessibility of PAs³⁵.

238 The type of fermentation also influences PA content. An alternative to the use of dry yeast is the use of
239 sourdough. Leavening with sourdough consists in the use of a starter, represented by a piece of dough from a
240 previous batch, which is fermented and stored under controlled conditions of temperature and humidity. The
241 intense acidification markedly influences the sensory and shelf-life features of the baked goods. With
242 sourdough, dough acidification and leavening capability is determined by the interactions between lactic acid
243 bacteria and yeasts. This kind of fermentation has a well-established role in improving flavor, structure, and
244 shelf-life of rye and wheat breads.

245 Sourdough fermentation has been demonstrated to increase the bioaccessibility of PAs as, for example,
246 Liukkonen et al. (2003)⁶⁴ found that this type of fermentation increased the content of methanol-extracted
247 phenolic compounds, in addition to demonstrating an increase in antioxidant capacity^{43,60}. As mentioned
248 above, low pH favors the hydrolysis of both esters and glycosides of PAs^{50,51}. However, different lactic acid
249 bacteria strains exhibit varying abilities in enhancing the extraction of free phenolics, with, for example, the

250 maximum increase in FA - in whole grain barley and oat groat when *Lactobacillus johnsonii* LA1,
251 *Lactobacillus reuteri* SD2112, and *Lactobacillus acidophilus* LA-5 were used⁶⁵.

252 Lastly, baking is considered the most important stage of the bread-making process. During baking, the
253 exchange of heat (as the dough heats up) and material (as the dough loses water/humidity) causes physical,
254 chemical and biochemical changes resulting in the transition from foam to sponge state and the
255 diversification between crust and crumb.

256 It is assumed that antioxidants, including PAs, contained in grains are lost during thermal treatments, due to
257 degradation, oxidative condensation, or decomposition of thermolabile phenolics caused by high
258 temperature^{66,67}. However, the most recent research has reported that baking increases the total PA and FA
259 levels^{35,59,68}, likely due to the intense heat that makes PAs more bioaccessible. Yu and Beta (2015)³⁵ found
260 higher contents of soluble FA and *p*-hydroxybenzoic acids in bread crumb compared to crust, suggesting that
261 some free PAs are thermally labile (since there is a higher, more intense heat in bread crust). However,
262 higher levels of insoluble PAs can be found in the crust^{35,59,68}. Heat stress could cause degradation of
263 conjugated polyphenolic compounds resulting in an increase in free PAs, which has been demonstrated in
264 wheat⁵³. This would improve bioavailability of phenolic compounds since it is believed that free PAs are
265 more readily available than bound PAs⁶⁹. The effect of baking temperature on free or bound PAs can vary
266 due to the nature and source of phenolic compounds as well as the baking method (e.g. yeast vs.
267 sourdough)^{51,70}.

268 Unlike temperature, baking time does not seem to affect total PA content of wholegrain bread, as
269 demonstrated in one study comparing breads baked at 10, 20 or 35 minutes⁶⁸. Additionally, Maillard reaction
270 that occurs during baking may contribute to the formation of new phenolic structures^{68,71}. Angioloni and
271 Collar (2011)⁷² demonstrated that some PAs, such as protocatechuic, syringic, SA and FA, were detected in
272 bread but not in the raw flour. This has also been shown in studies conducted with bread made from
273 pigmented wheat³⁵ and rye whole meal⁵⁰.

274 Furthermore, one study demonstrated that although there is a measurable decrease in total PA and FA
275 content that occurs during dough preparation, their concentrations significantly increased after baking to
276 levels that surpassed those measured prior to dough preparation⁵⁹. The baking process, however can have
277 different effects depending on the type of grain used. For example, in breads made with pseudocereals (e.g.

278 amaranth, quinoa, and buckwheat), polyphenol content has generally been found to be reduced in the final
279 bread product when compared to the original grains^{66,67}.

280

281 **BIOACCESSIBILITY OF PHENOLIC COMPOUNDS IN BREAD: *In vitro* studies**

282 **Methods of assessment of bioaccessibility: Static vs. Dynamic Methods**

283 Bioaccessibility is the determination of the amount of bioactive compounds potentially absorbable from the
284 gut lumen, and can be measured using different methods which simulate *in vivo* digestion. Several *in vitro*
285 methods have been developed to investigate the effect of the food matrix and of different processing
286 techniques on the ability of nutrients or bioactive compounds, like polyphenols, to become available to
287 absorption⁷³. These methods try to mimic *in vivo* digestion by simulating the oral, gastric and small intestinal
288 phases and, occasionally, large intestinal fermentation⁷⁴.

289 There are two general categories of methods: static and dynamic (non-static). In static models, products
290 remain largely immobile in a single bioreactor, and the ratios between meal, enzymes, salt, bile acids and all
291 other substrates of the biological digestive reactions are kept constant at each phase of digestion. Static
292 methods can differ in incubation time and characteristics of the digestive juices, namely the concentrations of
293 the enzymes resulting from the preparation, for example by the addition of specific enzymes to inorganic and
294 organic solutions. They can be also adjusted for pH on the basis of the specific gut compartment, as static
295 methods consist of multiple phases, including oral, gastric and intestinal, each of which can vary slightly in
296 different studies.

297 In the oral phase, the incubation time of the test sample can vary between 2 and 30 minutes^{74,75} with either: i)
298 human buffered saliva with phosphate or saline solution⁷⁴; ii) α -amylase solution^{75,76}; or iii) saliva solution
299 prepared with different salts and with the addition of α -amylase, uric acid and mucin⁷⁷. Some studies bypass
300 the oral phase^{72,78} possibly because a significant contribution to the digestive process is not expected in this
301 stage due to the short time during which food is in contact with saliva in *in vivo* conditions⁷⁹.

302 In the gastric phase, a pepsin solution is normally used and incubation time can vary between 1 and 2
303 hours^{72,74-78}. The addition of mucin has also been reported⁷⁷. Furthermore, hydrochloric acid is commonly
304 used to more accurately simulate *in vivo* gastric conditions⁷⁹.

305 In the intestinal phase, neutralization as well as incubation with pancreatic enzymes is set up. The enzymes
306 used in most studies include pancreatin⁷⁴, a bile/pancreatin solution^{72,76,78}, or a duodenal juice including
307 pancreatin, lipase and bile⁷⁷. Incubation time can vary from 2 to over 24 hours^{74,76,77}.

308 After gastrointestinal digestion simulation, the point at which bioaccessibility determination of compounds
309 of interest occurs can also vary. One method is to centrifuge or filtrate the sample mixture to measure the
310 bioaccessibility of compounds based on the levels present in the supernatant. An alternative method includes
311 the use of a dialysis membranes, which allows for the discrimination between high and low molecular weight
312 components³¹. When a dialysis tube is used, the undigested material (the fraction remaining inside the tube)
313 can be analysed for the content of the nutrient/bioactive compound under study (e.g. PAs) and then the
314 bioaccessibility can be obtained as a difference from that measured in the sample before digestion^{74,76}. The
315 time at which the dialysis tube is used may vary, as in some works it is used immediately after the gastric
316 phase⁷⁴ while in others after the intestinal digestion phase⁷⁶.

317 In general, static methods are quick, cost-effective and can be used to assess effects on several nutrients and
318 bioactive compounds resulting from changes made to the food matrix, by changing the raw materials or
319 processing techniques used, compared to the reference material or to the original food matrix.

320 The main limitations of static methods are that they do not provide the most accurate simulation of the
321 complex dynamic physiological processes occurring during *in vivo* conditions. This has led to the
322 development of dynamic (non-static) digestion models. A common and very sophisticated gut model to
323 simulate the human digestive system was developed by The Netherland Organization for Applied Scientific
324 Research⁸⁰. Their commercial gastrointestinal model, also known as the TIM system, is a multi-
325 compartmental dynamic computer-controlled model that has been successfully used to study the
326 bioaccessibility of many compounds including vitamins and minerals, as well as phenolics^{81,82}. The TIM
327 system simulates the dynamic conditions occurring in the four main gastrointestinal compartments: stomach,
328 duodenum, jejunum and ileum. All parameters, including gastric and small intestinal transit, flow rate,
329 composition of digestive fluids, temperature, pH, and removal of water and metabolites, are all remote-
330 computer controlled. In the jejunal and ileal compartments, a dialysis system allows for the removal of
331 digestion products, isolating the dialysate fraction, which contains the bioaccessible products from the
332 “unabsorbed” sample.

333 Overall, the use of realistic concentrations of digestive enzymes, pH levels, transit times appropriate to each
334 digestion step, and salt concentrations, among other factors, contribute to a more accurate simulation of the
335 gastrointestinal tract. Furthermore, the removal of the products of digestion and the appropriate mixing at
336 each stage of digestion in the use of dynamic methods may represent crucial points in mimicking
337 physiological conditions *in vivo*.

338 **In vitro studies investigating effects of altering raw materials on phenolic bioaccessibility**

339 Table 2 shows the main findings of all studies identified in the literature and evaluating the bioaccessibility
340 of phenolic compounds in bread. Among the different potential strategies to apply in the bread-making
341 process to increase the bioaccessibility of phenolic compounds in breads, as summarized in Table 1, the
342 efficacy of using different raw materials has been the most investigated. In particular, the majority of the
343 studies has explored the bioaccessibility in breads made by using different types of cereals or pseudocereals.
344 As expected, wheat-based breads (both white and wholegrain) were the most investigated (in all 9 studies),
345 with few investigating rye (2/9), oat (1/9) and barley (1/9), either alone or mixed. Among pseudocereals,
346 buckwheat breads were analysed in two out of the nine studies.

347 Almost all studies included wheat bread as an internal control to be compared with breads made with
348 different cereals. Generally, white bread is characterized by a low bioaccessibility of PAs, partially
349 ascribable to the very low FA content in the samples⁸³, especially in its free form. Three studies, investigated
350 the bioaccessibility of PAs in white breads following digestion by the dynamic TIM system, expressing
351 results as the percentage of PAs in the dialysate in relation to the original sample^{69,83,84}. In the first study by
352 Mateo Anson et al. (2009 a)⁶⁹, FA was undetected in the dialysate-samples, whereas in the second study,
353 conducted by the same authors⁸⁴, 4.9% FA bioaccessibility was reported. CA and SA were measured in the
354 second study⁸⁴, but they were not detected in the dialysate, post-digestion. The third study by Hemery and
355 colleagues (2010)⁸³ found a 10.2% FA bioaccessibility.

356 In another study conducted by Angioloni and Collar (2011)⁷², the authors found a 58% bioaccessibility of the
357 total phenolics (measured as Total Phenolic Content, TPC) in the supernatant from static *in vitro* digestion of
358 wheat bread. This is similar to a second study conducted by the same authors, in which they found ~84%
359 TPC bioaccessibility in wheat bread, although in this latter study the percentage bioaccessibility was
360 calculated from the initial TPC in flour as opposed to the bread, as is typically done⁷⁸.

361 The differences in bioaccessibility in white wheat bread found in the latter two studies (58% and 84% for
362 TPC) compared to the former three studies (0% FA, 4.9% FA and 10.2% FA) may be linked to the former
363 three measuring FA only, using chromatographic methods, while the latter two measured total phenolics
364 using the Folin-Ciocalteu method. In addition to potential differences due to type of *in vitro* method (*i.e.*
365 static vs. dynamic), further sources of variability might include the phenolic content in the raw materials, as
366 well as the state of the test samples used for post-digestion measurements (*i.e.* dialysate samples in former 3
367 versus supernatant and precipitate used in the latter 2 studies).

368 Three studies compared the bioaccessibility of phenolic compounds in white bread with respect to whole
369 wheat bread^{69,75,83}. As expected, the whole wheat breads had higher initial PA content, due to the
370 preservation of the outer layers of the kernels (e.g. 9-12-fold higher FA content in whole wheat versus white
371 bread). This contributed to a greater net content of bioaccessible PAs, demonstrating how the use of different
372 raw materials is a valid strategy for this purpose. However, although the net content of bioaccessible PAs in
373 whole wheat bread is higher than that of white bread, the bioaccessibility was higher in white breads
374 compared to whole wheat (e.g. 4.9% versus 1.1% in Mateo Anson et al. (2009b)⁸⁴ and 10.2% FA vs. 2.9%
375 FA in Hemery et al. (2010)⁸³). Nevertheless, other studies have observed much higher bioaccessibility for
376 specific PA in whole wheat breads. For example, Dall'Asta, *et al.* (2016)⁷⁴ found a 13.1% FA
377 bioaccessibility in whole grain bread, but this may be due to differences in the methods used, with this latter
378 study using a static digestion model. Variations in bioaccessibility in whole wheat breads may also differ due
379 to the type of whole wheat or whole grain bread used, as the former two studies produced breads from flour
380 at lab level, while the latter used a commercial whole grain bread which may have been exposed to different,
381 perhaps greater, degrees of processing. Furthermore, there seem to be no differences in bioaccessibility for
382 different types of PA. For example, FA appears to have lower percentage bioaccessibility compared to CA
383 and SA, regardless of the analytical method^{74,83}. This may be due to different distributions of phenolic
384 compounds in the free, conjugated, and bound forms.

385 Szawara-Nowak and colleagues (2016)⁷⁵, following *in vitro* digestion of white wheat bread, found a soluble
386 fraction of these compounds quite comparable to the content in dark wheat bread (~9 mg rutin equivalent/g
387 dry weight). They reported, for both white and dark bread, an exceptionally large increase in rutin post
388 versus pre-digestion (~20 and ~9 fold, respectively), which is much greater compared to any other study.

389 Similar unexpected increases following digestion were also found with increasing substitution of buckwheat
390 flour (both white and roasted) in white and dark breads⁷⁵. Authors hypothesized this may be due to an
391 increase in the extractability of phenolic compounds resulting from the parameters set in their *in vitro*
392 digestion, including pH, temperature, incubation times, and extraction solvent.

393 As reported in Table 1, a strategy to increase PA content in bread includes modifications to the raw
394 materials. The use of different types of cereals or pseudocereals, or a mixture of them, is a strategy to
395 increase PA content which has become increasingly common⁸⁵. In the above-mentioned study, Angioloni
396 and Collar (2011)⁷² assessed the differences in TPC in breads made with oat, rye, buckwheat and wheat
397 flours. Among the four breads made with 100% of one single type of flour, the TPC (measured by Folin-
398 Ciocalteu method) in the initial bread was highest in buckwheat (808 mg GAE/kg), followed by wheat (685
399 mg GAE/kg), oat (643 mg GAE/kg) and rye (536 mg GAE/kg). Following *in vitro* digestion, although the
400 bioaccessibility of TPC was greatest in the rye bread (62%), the net PA content was greatest in the 100%
401 wheat bread (401 mg GAE/kg) with 58% bioaccessibility, followed by buckwheat (366 mg GAE/kg; 45%),
402 rye (334 mg GAE/kg; 62%) and then oat (264 mg GAE/kg; 41%). The lower bioaccessibility in the
403 buckwheat and oat breads may be due to their substantially greater fibre and protein contents in the
404 respective flours (13.8% and 17.4%, and 18.9% and 21.5%, respectively) compared to the white wheat and
405 rye breads (2.2% and 12.6%, and 14.6% and 9.6%, respectively). The higher fibre and protein content may
406 partially prevent the digestive enzymes to free bound PAs, thus limiting their bioaccessibility. The same
407 study also assessed blends of flours, specifically the multigrain bread “blend 15%” (oat:rye:buckwheat:wheat
408 15:15:15:55, “blend 20%” (20:20:20:40) and “blend 25%” (25:25:25:25), where the TPC in the initial bread
409 increased with the increase in wheat flour replacement (from 592 to 745 to 916 mg GAE/kg). Interestingly,
410 the higher the substitution level of wheat flour by minor cereal and pseudocereal, the lower the percentage of
411 TPC bioaccessibility, with the highest value (80%) reached with the 15% blend. However, the net TPC was
412 comparable between the 3 blends (472, 549, 504mg GAE/kg corresponding to the 15, 20, 25% blends), and
413 was actually greater than any of the 100% breads (401, 366, 334 and 264 mg GAE/kg, for wheat, buckwheat,
414 rye and oat breads, respectively). Therefore, there may be some influential effect on PA bioaccessibility
415 resulting from mixed grains, regardless of the actual quantities of each individual type. Comparing the 100%
416 wheat flour bread with the 15% blend, which was 55% wheat flour, it is interesting to notice that that the

417 initial TPC content of the 100% wheat flour bread was higher (685 vs. 592mg GAE/kg TPC), yet the final
418 bread TPC is higher in the 15% blend (472 vs. 401mg GAE/kg).

419 In another study by the same authors, a 40% barley bread (made by replacing 40% wheat flour with barley
420 flour) showed no difference in net TPC (597 vs. 598 mg/100g, respectively) and a much lower %
421 bioaccessibility (60% vs. 84%, respectively, although the difference in TPC in the flour was higher (1003
422 mg/100g vs. 713 mg/100g)⁷⁸. Perhaps the specific barley flour used, commercial barley flour, had low
423 bioaccessibility due to greater fibre content (4.01 vs. 1.15 g/100g, in the respective breads). When the type of
424 barley flour was changed to a high β -glucan barley flour, the percentage bioaccessibility was still lower
425 compared to the bread made from refined common wheat flour (42% vs. ~84%, respectively), again likely
426 due to the higher fibre content (11.91 vs. 1.15 g/100g in the respective breads). However, the net TPC was
427 much greater in the high β -glucan barley bread compared to the 100% white wheat bread (857 vs. 598
428 mg/100g, respectively) because the TPC in the raw flour was ~3-fold higher (2197 vs. 713 mg/100g,
429 respectively). Beta-glucan is a soluble, viscous-type fibre, which may therefore contribute to the low PA
430 bioaccessibility since β -glucans can produce viscous gels able to entrap nutrients and phytochemicals,
431 including phenolics, as previously hypothesized⁷². This may also explain part of the particularly lower
432 bioaccessibility in the oat bread found in the study discussed above⁷², since oats are also a rich source of β -
433 glucan soluble fibre. Overall, these studies demonstrate that the types of grain flour used in blends may be
434 influential on PA bioaccessibility.

435 Another way to increase PA content in breads by modifications of raw materials includes the addition of
436 selected fractions from the original grain. One of the most commonly used fractions is the cereal bran, as it is
437 a recognized source of phenolics, including PAs. Mateo Anson *et al.* (2009b)⁸⁴ compared a wholemeal bread
438 to a wholemeal bread added with native wheat bran. Although they found the same FA bioaccessibility in
439 both breads (1.1%), the net FA content in the wholemeal bread plus bran was greater, since the bread plus
440 bran had a greater initial content of FA (1300 μ g/g vs. 800 μ g/g). The potential reason why the FA
441 bioaccessibility was the same between the breads is because the bioaccessibility of FA is mainly associated
442 with the amount of free FA present in breads, and the FA in the bran is mostly bound. Mateo Anson *et al.*
443 (2009b)⁸⁴ demonstrated a strong correlation between the amount of free FA and bioaccessibility among five
444 breads. This hypothesis is further supported by the study of Koistinen *et al.* (2017)⁷⁶, where the authors

445 compared wheat bread made with bioprocessed rye bran to the same bread made with native rye bran and
446 found that FA bioaccessibility was significantly greater in the bread with the bioprocessed rye bran (88% vs.
447 51%, respectively). This was also reflected in the bioaccessibility of total PAs (89% vs. 53%, respectively).
448 The bioaccessibility was not directly calculated in this study. However, by calculating it as the difference
449 between polyphenol content in the original sample and the residue of the enzymatic digestion⁸⁶, percentage
450 bioaccessibility of PAs was inferred.

451 In addition to the use of bran, bread can be enriched with the polyphenol-rich aleurone fraction, as was
452 investigated in two studies^{69,74}. In the study by Mateo Anson *et al.* (2009a)⁶⁹, the addition of aleurone
453 resulted in a substantial increase in initial FA in the bread compared to the white bread (2290 µg/g and 33.5
454 µg/g, respectively). After *in vitro* dynamic digestion, an increase in FA bioaccessibility was detected in the
455 aleurone-enriched bread (0.57%) compared to white bread (not detected). Furthermore, Mateo Anson *et al.*
456 (2009a)⁶⁹ demonstrated that the aleurone-enriched bread had a level of FA bioaccessibility that was ~60%
457 lower than that found in a raw flour which had free FAs added (which was used as a “positive” control). In
458 the aleurone-enriched bread, the majority of FA was present in bound form and only 20 µg/g as free FAs.
459 Considering that only free and conjugated phenolic compounds are readily available for absorption, these
460 results further support the consideration that free phenolic compounds are the major contributors to the
461 bioaccessibility of PAs. Conversely, bound phenolics, being largely attached to undigested cell wall
462 polysaccharides, are mainly retained into the material reaching the colon.

463 The static model study by Dall’Asta *et al.* (2016)⁷⁴ showed instead that aleurone-enriched bread resulted in
464 bioaccessibility values 2.5-fold to 4.4-fold greater compared to whole grain bread for various PAs, including
465 a 3-fold greater bioaccessibility for FA. These results are particularly interesting, since the aleurone bread
466 had approximately half the amount of PAs compared to the wholegrain bread (total FA 70.67 vs. 144.78
467 mg/100g, respectively). Although the results from this latter study contrast the ones of Mateo Anson *et al.*
468 (2009a)⁶⁹, they are supported by a previous study where it was reported that, in addition to the free form, a
469 relevant percentage of the bound fraction may become available for absorption following digestion⁸⁷. The
470 mechanisms through which aleurone additions may influence PA bioaccessibility in the two studies may be
471 ascribable to several factors. In addition to the differences between the *in vitro* method used (TIM versus
472 static), the studies differed in the applied digestion length (6 versus 24 hours), in the aleurone content (22%

473 vs. 9.3% aleurone flour in the final dough), and in the kind of phenolic compounds considered (*i.e.* the
474 consideration of di- and tri-FA in the work of Dall'Asta *et al.* (2016)⁷⁴). Regardless, both studies demonstrate
475 that the use of the polyphenol-rich aleurone fraction may represent a valuable source of phenolics and as an
476 attractive strategy for producing breads with bioaccessible PAs, along with the advantage of more acceptable
477 sensory characteristics.

478 **In vitro bioaccessibility studies investigating effects of pre-processing techniques in bread-making on**
479 **polyphenolic content**

480 Beyond using different raw materials to influence PA content, innovative technologies have been developed,
481 including pre-processing techniques, with the aim to improve the release of bound phenolic compounds and
482 thus their bioaccessibility. Biotechnological processing and dry-fractionation of wheat bran are two types of
483 technologies that have thus far been investigated in *in vitro* digestion studies assessing bioaccessibility of
484 phenolic compounds in bread^{76,83,84}.

485 Fermentation and enzymatic treatment are two biotechnological processing techniques applied during bread-
486 making, which have been investigated on their effect on the bioaccessibility of FA, CA and SA. One study
487 compared a wholemeal bread with native wheat bran to one where the wheat bran had been fermented and to
488 another where the wheat bran had been both fermented and enzymatically treated with xylanase, β -
489 glucanase, α -amylase, cellulase and ferulic acid esterase⁸⁴. All three breads had the same initial content of
490 FA, CA and SA. However, after a dynamic digestion method was applied, the bioaccessibility of FA was
491 twice as high in the bread with fermented wheat bran and 5-fold higher in the bread with fermented and
492 enzymatically treated wheat bran, compared to the bread with native wheat bran. A slightly smaller but
493 similar trend was observed for CA and SA. The great increase in bioaccessibility in the bread with
494 bioprocessed bran may be due to the hydrolysis of different wheat fibre polymers resulting from to the
495 hydrolytic enzymes, which may lead to a structural breakdown of bran cell walls.

496 Mandak and Nystrom (2013)⁷⁷ also evaluated the effect of enzymatic treatment, and assessed the
497 bioaccessibility of steryl ferulates, which are phytosterols that can be esterified to FA, in breads made with
498 two types of wheat flour, either with or without the use of the enzymes cellulase or xylanase, alone or in
499 combination. The bioaccessibility of steryl ferulate (calculated as the percentage in the supernatant compared
500 to the total extractable amount) was generally very low (0.01-0.25%), although when both enzymes were

501 used, bioaccessibility increased from 0.01 to 0.25% in wholegrain breads, but only from 0.09 to 0.10% in
502 baking flour breads. The differences in effect of enzymatic treatment seen in this study versus the study by
503 Mateo Anson *et al.* (2009b)⁸⁴ may be: i) the specificity in the phenolic compounds assessed (steryl ferulates
504 vs. PAs); ii) the specific enzymes used and the number and combination of them (xylanase and cellulase vs.
505 β -glucanase, xylanase, α -amylase and ferulic acid esterase); iii) the method of bread preparation (direct
506 incorporation of the enzymes to the flour vs. preliminary bioprocessing of bran); and iv) the digestion
507 method employed (static vs. dynamic).

508 As previously mentioned, Koistinen *et al.* (2017)⁷⁶ recently investigated the bioaccessibility of phenolic
509 compounds in a bioprocessed (by enzymatic treatment and fermentation) rye bran added to wheat bread, and
510 found a stunning 88% bioaccessibility of FAs. Bioaccessibility was therefore much higher than that of the
511 two previous studies, possibly because a considerable amount of phenolic bound compounds became
512 available due to the addition of enzymes and the activation of endogenous enzymes resulting from
513 fermentation.

514 The bioaccessibility of PAs in bread was also increased when wheat bran was dry-fractionated. Hemery *et al.*
515 (2010)⁸³ analysed free, conjugated, bound and total FA, SA and CA in bread made following bran ultra-fine
516 grinding and bran electrostatic separation. They found that the finer the bran particles in bran-rich breads, the
517 more bioaccessible the PAs (following Tiny-TIM digestion), with a very strong correlation between FA
518 bioaccessibility and the proportion of small particles (10-20 μ m diameter). The bioaccessibility of SA was
519 generally much higher than that of CA or FA (26-33% versus 6-13% and 2.5-3.4%), likely because SA is
520 mainly present in the conjugated form and within the aleurone grains⁸⁸. Furthermore, although the breaking
521 of covalent bonds during extensive milling contributes to increased bioaccessibility⁸⁹, the particle size of the
522 samples seems to play a role in determining the bioaccessibility of phenolic compounds, possibly through an
523 improvement of the extractability resulting from micronization⁹⁰. The described study also found SA
524 bioaccessibility was highly correlated to the proportion of small particles (<10 μ m diameter), and the authors
525 furthermore evaluated also bread made with positive and negative fractions obtained by electrostatic
526 separation of bran, after the highest level of grinding (cryo-ultrafine), and demonstrated these to have the
527 highest amount of bioaccessible PAs. The charge of these particles was influenced by the type of cell walls
528 (branched and cross-linked vs. linear oligosaccharides), with separation between fibre-rich particles of

529 pericarp (outer cell wall), rich in highly branched and cross-linked arabinoxylans (negatively charged) and
530 particles rich in β -glucan, FA and CA from aleurone cell walls (positively charged)⁹¹. These results provide
531 insights for the improvement of electrostatic separation processes able to select specific fractions rich in free
532 and conjugated PAs⁴⁰.

533 Overall, the studies investigating the bioaccessibility of phenolic compounds in bread suggest alterations,
534 such as the incorporation of polyphenol-rich raw materials and, especially, the application of different bio-
535 processing techniques represent promising strategies to increase the amount of bioaccessible phenolic
536 compounds in bread. The significant variations among the *in vitro* methods used impede a proper
537 comparison of the results across studies and make the possibility to deduce general findings very difficult. To
538 circumvent this, Minekus *et al.* 2014⁹² recently published an international consensus paper aimed at
539 introducing a standardised *in vitro* digestion method to analyse food, providing recommendations for every
540 step of digestion. Adoption of this standardized method will assist in comparison of multiple study results in
541 the future, allowing for clearer conclusions to be drawn.

542

543 **BIOAVAILABILITY OF PHENOLIC COMPOUNDS IN BREAD: *in vivo* studies**

544 Determining the content of bioactive compounds in food products or their sole bioaccessibility *in vitro* is not
545 sufficient per se to predict their potential health effects *in vivo*. Therefore, *in vivo* studies are important to
546 determine the bioavailability of PAs in order to understand the amount of PA actually absorbed post-
547 ingestion, becoming therefore available to elicit health effects.

548 A review of the literature identified 5 studies investigating the bioavailability of PAs from standard versus
549 bioprocessed bread (Table 3). The most common methodology used *in vivo* to assess phenolic bioavailability
550 is represented by acute studies, where subjects are provided a single-dose of the test food and biological
551 samples (e.g. blood, urine) are collected pre- and post-consumption. The changes, therefore, reflect the
552 ability to absorb polyphenols from a complex food matrix⁹³. Three out of the five identified studies were
553 single-dose acute studies, with 2 evaluating the bioavailability of phenolics in bread in urine and plasma^{49,94}
554 and 1 in urine alone⁹⁵.

555 Bioavailability was calculated in all studies as the ratio between the amount of the excreted phenolic
556 compounds and the amount provided with in the fed bread sample. Bresciani *et al.* (2016)⁹⁴ specifically

557 detected and quantified secondary metabolites of phenolic compounds and described the bioavailability as
558 the sum of these conjugated metabolites, while Lappi et al. (2013)⁹⁵ and Mateo Anson et al. (2011)⁴⁹
559 performed an enzymatic hydrolysis of the urinary sample by using a mixture of β -glucuronidase and
560 sulfatase from *Helix pomatia*. This reaction allows to cleave the glucuronic and sulfonic moieties of the
561 phase II metabolites and to detect the only aglycones, to which the bioavailability is accounted for.

562 As discussed, raw materials as well as bioprocessing techniques in bread-making play important roles in the
563 bioavailability of phenolic compounds in breads. Product innovation in these acute studies was based on
564 three main strategies: i) the addition of aleurone fraction to commercial wheat breads⁹⁴; ii) bioprocessing of
565 wheat bran added to a whole grain bread⁴⁹; and iii) the use of rye bread and rye bran⁹⁵.

566 All 3 acute studies evaluated the urinary bioavailability of FA. Bresciani et al. (2016)⁹⁴ fed healthy
567 volunteers, on three separate days, a wholegrain bread and a 6% w/w aleurone-enriched bread at two
568 different servings of 94 g and 190 g, containing 43 mg and 87 mg total FA, respectively. Results showed a
569 significant 2-fold higher FA bioavailability (as the sum of FA metabolites ferulic acid-4'-O-sulfate,
570 dihydroferulic acid-4'-O-sulfate, and dihydroferulic acid-O-glucuronide) in urine of volunteers fed with the
571 single portion of the aleurone-enriched bread compared to wholegrain bread and to the double portion of
572 aleurone-enriched bread. Intriguingly, no significant difference was found in urinary FA bioavailability
573 between the double portion of aleurone-enriched and wholegrain breads (~5% and ~4%, respectively). The
574 authors commented that the higher bioavailability derived from the lower ferulic consumption in the single
575 compared to double portion of aleurone-enriched bread may be due by a reduction in the capacity to
576 metabolize and absorb PAs as intake increases.

577 Mateo Anson et al. (2011)⁴⁹ demonstrated similar results when breads were standardized to contain the same
578 initial total PA amount. Specifically, they found 10% FA bioavailability in the bread with bioprocessed bran
579 compared to 4% in the whole wheat control bread with native bran (21.34 mg/24h vs. 9.89 mg/24h FA in
580 urine, $p < 0.05$). Furthermore, Lappi et al. (2013)⁹⁵ found a 2.5-fold greater urinary FA excretion after
581 consumption of whole wheat bread with bioprocessed rye bran compared to the same whole wheat bread
582 with native rye bran and with control wheat. For a thorough comprehension of the results of this study, it is
583 important to consider the initial amount of FA in the fed bread. Indeed, the control wheat bread in this study
584 showed a 3.2% FA bioavailability, as per excretion in urine, even if the initial FA intake was much lower

585 compared to both the rye or the bioprocessed rye brans. Thus, the total FA urinary excretion was lower (0.27
586 mg/d in control whole wheat bread vs. 1.66 mg/d from bioprocessed rye bran bread vs. 0.45 mg/d in native
587 rye bran bread, corresponding to 3.2%, 1% and 0.4% FA bioavailability, respectively). Therefore, although
588 the percentage bioavailability may be higher, if the initial intake is lower, the total amount absorbed may
589 nevertheless be lower.

590 The application of bioprocessing techniques to breads, similarly, elicited increased bioavailability for SA and
591 other PAs. The study by Mateo Anson and colleagues (2011)⁴⁹ found that the amount of SA in 24-hour urine
592 corresponded to a 15% and 7% bioavailability in bioprocessed bran and control breads, respectively.
593 However, the bioavailability for CA equalled 2% for both the bioprocessed and control breads. Lappi *et al.*
594 (2013)⁹⁵ showed a 0.6% SA bioavailability for the bioprocessed bread compared to a 2.8% for white wheat
595 bread, and generally all three breads were characterised by a ~4-fold lower SA bioavailability compared to
596 the white wheat bread. In spite of this, the bioprocessed bread showed the highest excreted SA net amount
597 (0.23 mg vs. 0.06-0.12 mg in the other three breads). Similar results were found for CA bioavailability.
598 Intriguingly, Mateo Anson *et al.* (2011)⁴⁹ evaluated the percentage vanillic acid bioavailability based on 24-
599 hour urine excretion and demonstrated 160% and 104% bioavailability in the bioprocessed and control
600 breads, respectively, and both had similar initial concentrations in breads (0.018 mg/g and 0.017 mg/g,
601 respectively), thus the bioprocessed bran bread resulted in greater vanillic acid absorption. Authors did not
602 provide a possible explanation for such high recoveries, which could be at least partially attributable to an
603 insufficient initial extraction of phenolics from the bread.

604 Two studies also evaluated blood concentrations of phenolic compounds after bread consumption^{49,94} and
605 both demonstrated increased hippuric and hydroxyhippuric acid plasma levels after bread consumption.
606 However, being degradation products from several different metabolic pathways, these two catabolites
607 cannot be considered uniquely associated to polyphenol metabolism²². The second most concentrated
608 polyphenol compound in plasma was FA, together with its main phase II conjugates. Bresciani *et al.* (2016)⁹⁴
609 found concentrations of the main FA metabolites (ferulic acid- 4'-O-sulfate and dihydroferulic acid-4'-O-
610 sulfate) ranging from 66 to 100 nmol/L at 90 minutes after bread intake, with no significant differences
611 among the various breads. Mateo Anson *et al.* (2011)⁴⁹, however, found a significantly higher plasma FA
612 concentration from bioprocessed bread (2.7 $\mu\text{mol/L}$) compared to control bread (0.9 $\mu\text{mol/L}$). The contrast of

613 these results may be explained by the different initial intakes of FA, as, although the concentration of FA in
614 breads were similar, in the latter study⁴⁹ the subjects consumed 3 times as much bread (300 g vs. 94 g) and
615 thus had a 3-fold higher FA intake. Two studies investigated the consumption of rye bran breads in the
616 context of a dietary intervention. The study by Harder *et al.* (2004)⁹⁶ compared 250 g/d of rye bran-enriched
617 products with 250 g/d of control wheat products (Vitacell[®]) consumed for 6 weeks in a randomized,
618 crossover designed intervention with a 4-week washout in 18 healthy postmenopausal women. Juntunen *et*
619 *al.* (2000)⁹⁷ similarly compared the consumption of 4-5 slices/d of rye bread with wheat bread for 4 weeks in
620 a randomized, crossover design with a 4-week washout in 43 healthy volunteers (Table 3). Although it was
621 not possible to calculate the bioavailability of phenolic compounds in the breads because measurements
622 would have had to include phenolics⁴⁹ found in foods consumed during the rest of the daily diet,
623 measurements of phenolic metabolites in blood (plasma) and urine samples were compared after
624 consumption of the different bread interventions. Moreover, in the study by Harder *et al.* (2004)⁹⁶, in
625 addition to rye bread, the authors also included rye-enriched muffins and crisp bread products, thus making
626 the FA amount found in biological samples not originating solely from bread. These authors measured FA
627 concentration in 48-hour urine collections and found urinary FA excretion was ~2 mg/24hour for the habitual
628 diet (*i.e.* at baseline) and at the end of 6-weeks after the incorporation of white wheat bread (Vitacell[®]).
629 However, at the end of 6-weeks of the intervention with rye bran-enriched bread products, FA excretion was
630 2.5-fold higher ($p < 0.05$) compared with both the control wheat bread intervention (40.2% higher, $p = 0.001$)
631 and the baseline diet (39.8% higher, $p = 0.002$). Considering the 10.2 mg FA/day intake during the rye bran
632 intervention, the study demonstrated a recovery of 28% of FA metabolites.

633 Juntunen *et al.* (2000)⁹⁷ considered the plant lignans, secoisolariciresinol (SECO) and matairesinol (MAT),
634 which are found in large quantities in rye cereal-based products and bio-transformed by gut microbiota into
635 enterodiols and enterolactone (ENL), respectively, and the latter finally oxidized to ENL. After a 4-week
636 dietary intervention on either wheat or rye bread consumption, total ENL excretion in 24-hour urine samples
637 almost doubled after rye bread consumption (6.8 $\mu\text{mol/day}$ for men and 7.8 $\mu\text{mol/day}$ for women) compared
638 to the period with wheat control bread (4.0 $\mu\text{mol/day}$ for men and 3.7 $\mu\text{mol/day}$ for women). However, 24-
639 hour urine ENL concentration at the end of the rye intervention was not significantly different from the
640 baseline. Furthermore, there was no correlation between the intake of rye bread or plant lignans and ENL

641 urinary excretion, which is interesting considering the intake of rye bread was more than double during the
642 rye bread intervention compared to the habitual diet. Additionally, no difference in serum ENL
643 concentrations between pre- and post-rye intervention was observed and again, there was no correlation
644 between rye intake and serum ENL concentration. It is possible that a plateau of ENL is physiologically
645 reached independently from the intake of rye bread.

646 **CONCLUSIONS AND REMARKS**

647 Phenolic compounds are recognized for several beneficial effects on human health. These effects depend not
648 only on their content in food products but also on their ability to be absorbed and become available within
649 the human body. For this reason, *in vitro* and *in vivo* studies have been performed with the aim of
650 investigating the bioaccessibility and bioavailability of phenolic compounds, respectively, suggesting that the
651 use of specific raw materials (e.g. cereals/pseudocereals as alternatives to wheat, or specific cereal fractions)
652 or of pre-processing techniques might represent valuable strategies for enhancing the phenolic content in the
653 raw materials and for increasing the amount of bioaccessible and bioavailable compounds.

654 Unequivocal conclusions could not be drawn at present, as the available studies widely differ for fed
655 amounts of phenolic compounds and, more importantly, for the methodologies applied. This highlights a
656 great need for standardization of methodologies used in *in vitro* studies in order to be able to compare results
657 and draw conclusions on the potential usefulness of the application of innovative techniques to improve
658 phenolic bioaccessibility. The few *in vivo* studies identified also highlight the need for further research to be
659 carried out in this area to assess the effectiveness of the application of new strategies in the bread-making
660 process on phenolic bioavailability. With the ultimate goal of eliciting health benefits, intervention trials will
661 be required to assess if strategies that demonstrate effectiveness at increasing phenolics bioavailability
662 translate then to improvements in health outcomes in humans.

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665 The authors declare no conflict of interest.

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Table. 1 Potential strategies to increase bioaccessibility and bioavailability of phenolic compounds in bread

STRATEGY			REASON/MECHANISM	References
Raw materials	Type of grain/cereal	Whole grains	Keeping all the anatomic parts of the kernel, where phenolic compounds are located	Hemery <i>et al.</i> (2007) ⁴⁰
		Rye, Barley	Raw material naturally rich in phenolic compounds	Dykes and Rooney (2007) ⁹⁸
		Minor cereals		Taylor and Duodu (2015) ³²
		Pseudocereals		Alvarez-Jubete <i>et al.</i> (2010a) ⁶⁷
		Ancient grains		Abdel-Aal and Rabalski (2008) ³⁶
		Pigmented grains		Abdel-Aal <i>et al.</i> (2012) ³³ Abdel-Aal <i>et al.</i> (2016) ³⁴ Yu and Beta (2015) ³⁵
	Selected fractions	Bran	Anatomic parts of the kernel, rich in phenolic compounds	Rosa-Sibakov <i>et al.</i> (2015) ⁹⁹
		Aleurone layer		
Pre-processing	Fractionation	De-branning	Selection of phenolic-rich fractions	Blandino <i>et al.</i> (2013) ⁴¹ Martini <i>et al.</i> (2015) ⁴³ Zanoletti <i>et al.</i> (2017) ⁴²
	Physical treatment	Air classification	Selection of phenolic-rich layers	Verardo <i>et al.</i> (2011) ⁴⁴
	Mechanical treatment	Micronization	Ultrafine grinding which damages the fiber matrix and increases the phenolic compounds available for extraction	Zhu <i>et al.</i> (2010) ⁴⁵
	Bio-technological processes	Germination	Metabolic changes and/or increase in extractability by the activation of endogenous enzymes which break the bonds of bound phenolic compounds	Hubner and Arendt (2013) ⁴⁶ Alvarez-Jubete <i>et al.</i> (2010b) ⁶⁷
		Fermentation/leavening	Release of insoluble bound phenolic compounds by activity of exogenous enzymes	Katina <i>et al.</i> (2007) ⁴⁷ Zhang <i>et al.</i> (2014) ⁴⁸ Poutanen <i>et al.</i> (2009) ¹⁰⁰

		Enzymatic treatment	Addition of enzymes which act to increase free phenolic compounds available for extraction	Sørensen <i>et al.</i> (2003) ⁵² Moore <i>et al.</i> (2006) ⁵³
Bread-making process	Mixing and kneading		Release of bound phenolic compounds into free forms by mechanical action and/or activation of oxygenase and peroxidase	Hilhorst <i>et al.</i> (1999) ⁶² Abdel-Aal and Rabalski (2013) ⁵⁸
	Fermentation/ Leavening	Length of fermentation	Prolonged fermentation time increase the phenolic compounds available for extraction	Yu and Beta (2015) ³⁵
		Type of fermentation (sourdough vs dry yeast)	Increase in the release of insoluble bound phenolic compounds during sourdough fermentation favoured by the lowering of pH	Boskov Hansen <i>et al.</i> (2002) ⁵⁰ Konopka <i>et al.</i> (2014) ⁵¹
	Baking	Temperature	Possible decrease in phenolic content due to degradation (thermal labile) Possible increase in phenolic bioaccessibility due to the release resulting from intense heat E.g. The upper crust, exposed to the greatest heat, generally has the highest level of phenolic compounds	Vogrincic <i>et al.</i> (2010) ⁶⁶ Alvarez-Jubete <i>et al.</i> (2010) ⁶⁷ Lu <i>et al.</i> (2014) ⁵⁹ Gélinas and McKinnon (2006) ⁶⁸ You and Beta (2015) ³⁵
		Maillard Reactions	May result in newly generated phenolic compounds	Gélinas and McKinnon (2006) ⁶⁸ Michalska <i>et al.</i> (2008) ⁷¹
		Time	No known effect	Gélinas and McKinnon (2006) ⁶⁸

Table 2: Summary of in vitro studies investigating the bioaccessibility of PAs resulting from the alterations to the bread-making process

Reference (Method)	Type of grain	Type of bread	Phenols analyzed	Initial phenolic content†		Main findings	
				Total	Free	Net Content	% bioaccessibility
Mateo Anson <i>et al.</i> (2009) ⁶⁹ (Dynamic- TIM)	Wheat	a) White bread b) Aleurone-enriched bread (50% flour replacement; 22% in final dough)	- FA	µg/g FA a) 33.5 b) 2290	µg/g FA a) 2.4 b) 20	a) not detectable b) 0.69 mg free	a) not detectable b) 0.57% free
Angioloni and Collar (2011) ⁷² (Static)	Wheat, oat, rye, buckwheat	a) White wheat bread b) Buckwheat bread c) Rye bread d) Oat bread e) Blend 15%, Multigrain bread (oat:rye:buckwheat: wheat 15:15:15:55): f) Blend 20%, Multigrain bread (oat:rye:buckwheat: wheat 20:20:20:40): g) Blend 25%, Multigrain bread (oat:rye:buckwheat: wheat 25:25:25:25)	- TPC	mg GAE/kg a) 685 b) 808 c) 536 d) 643 e) 592 f) 745 g) 916	n/a	GAE mg/kg a) 401 b) 366 c) 334 d) 264 e) 472 f) 549 g) 504	a) 58% b) 45% c) 62% d) 41% e) 80% f) 74% g) 55%
Collar and Angioloni (2014) ⁷⁸ (Static)	Wheat, barley	a) White wheat bread b) 40% barley bread (40% wheat replaced with commercial barley flour) c) 40% high beta-glucan barley bread (40% wheat replaced with high β-glucan barley flour)	- TPC	mg/100g dw in flour a) 713 b) 1003 c) 2197	n/a	mg/100g bread as is a) 598 b) 597 c) 857	a) ~84%* b) ~60%* c) 42%* *based on initial content in flours
Szawara-Nowak <i>et al.</i> (2016) ⁷⁵ (Static)	Wheat, buckwheat	a) White wheat bread b) Dark wheat bread c) White wheat bread with white buckwheat flour	- TPC	a) 0.38 mg rutin eq./g dw b) 1.8 mg rutin eq./g dw	n/a	Soluble fraction: a) ~9 mg rutin eq./g dw b) ~9 mg rutin eq./g dw	(Extrapolated) a) ~20 folds b) ~5 folds

		(substitution from 10% to 50%) d) White wheat bread with white roasted buckwheat groats (substitution from 10% to 50%) e) Dark wheat bread with white buckwheat flour (substitution from 10% to 50%) f) Dark wheat bread with white roasted buckwheat groats (substitution from 10% to 50%)		c) ~8-fold increase with 50% substitution as compared to a) d) ~11-fold increase with 50% substitution as compared to a) e) f) increase from 1.5 to 3 times as compared to b)		c) up to ~11 mg rutin eq./g dw (50% substitution) d) up to ~12.5 mg rutin eq./g dw (50% substitution) e) up to ~10.5 mg rutin eq./g dw (50% substitution) f) up to ~10.5 mg rutin eq./g dw (50% substitution)	c) ~3.5 to ~6 folds d) ~3 to ~5 folds e) ~3 to ~6 folds f) ~2.5 to ~4.5 folds
Dall'Asta <i>et al.</i> (2016) ⁷⁴ (Static)	Wheat	a) Whole grain bread (commercial) b) Aleurone-enriched bread (commercial)	- FA - CA - SA - CFA	mg/100g dw a) 144.78 FA 1.51 CA 3.08 SA 0.83 CFA b) 70.67 FA 0.87 CA 3.96 SA 0.28 CFA	mg/100g dw a) 0.71 FA 0.02 CA ND SA 0.04 CFA b) 0.41 FA 0.05 CA 0.12 SA 0.02 CFA	mg/100g dw (calculated) a) 18.97 FA 0.15 CA 0.99 SA 0.16 CFA b) 28.76 FA 0.26 CA 3.15 SA 0.23 CFA	a) 13.1% FA 10.1% CA 32.2% SA 19.2% CFA b) 40.7% FA 29.5% CA 79.5% SA 83.3% CFA
Mateo Anson <i>et al.</i> (2009) ⁸⁴ (Dynamic, TIM)	Wheat	a) White bread b) Whole-meal bread c) Whole-meal bread with native wheat bran d) Whole-meal bread with	- FA - CA - SA	µg/g dw a) 86 FA 2 CA 9 SA b) 810 FA 20 CA 70 SA c) 1300 FA 40 CA 130 SA d) 1300 FA	µg/g free a) 3.6 FA 0.8 CA 0.9 SA b) 13 FA 0.9 CA 3.5 SA c) 12 FA 1.2 CA 4.6 SA d) 42 FA	µg/g (calculated) a) 4.2 FA n/a CA n/a SA b) 8.91 FA n/a CA n/a SA c) 14.3 FA 2.08 CA 2.73 SA	a) 4.9% FA n/a CA n/a SA b) 1.1% FA n/a CA n/a SA c) 1.1% FA 5.2% CA 2.1% SA d) 2.2% FA

		fermented wheat bran e) Whole-meal bread with fermented and enzymatic treated bran		40 CA 130 SA e) 1300 FA 40 CA 130 SA	1.5 CA 9.6 SA e) 100 FA 3.0 CA 9.9 SA	d) 28.6 FA n/a CA n/a SA e) 71.5 FA 3.96 CA 6.5 SA	n/a CA n/a SA e) 5.5% FA 9.9% CA 5.0 % SA
Hemery <i>et al.</i> (2010) ⁸³ (Dynamic, Tiny-TIM)	Wheat	a) White bread b) Whole bread (100% wheat grain) c) “Amb, medium” d) “Amb, fine” e) “Amb, ultrafine” f) “Cyro, ultrafine” g) “FES positive” h) “FES middle” i) “FES negative” <i>* coarse bran increasingly processed from c-f; FES=cryo particles separated by charge; middle is mixed</i>	- FA - CA - SA	µg/g dw a) 62.6 FA 2.5 CA 3.2 SA b) 793.2 FA 23.5 CA 40.7 SA c) 865.4 FA 24.7 CA 44.7 SA d) 898.5 FA 26.6 CA 41.0 SA e) 899.4 FA 26.8 CA 42.5 SA f) 869.8 FA 25.6 CA 44.2 SA g) 1072.7 FA 30.9 CA 41.3 SA h) 763.8 FA 22.0 CA 48.1 SA i) 625.8 FA 25.9 CA 38.1 SA	µg/g dw a) 1.2 FA 0.12 CA 0.09 SA b) 8.2 FA 0.28 CA 1.68 SA c) 12.4 FA 0.48 CA 1.17 SA d) 14.6 FA 0.55 CA 1.21 SA e) 15.6 FA 0.53 CA 1.17 SA f) 16.4 FA 0.56 CA 1.21 SA g) 12.4 FA 0.47 CA 1.36 SA h) 17.9 FA 0.65 CA 1.01 SA i) 15.5 FA 0.56 CA 1.28 SA	µg/g dw a) 6.4 FA 0.87 CA 3.3 SA b) 22.7 FA 1.38 CA 18.1 SA c) 21.7 FA 1.49 CA 12.2 SA d) 26.2 FA 1.83 CA 13.8 SA e) 30.7 FA 2.47 CA 15.4 SA f) 26.7 FA 3.32 CA 11.7 SA g) 31.8 FA 3.59 CA 1.67 SA h) 23.0 FA 3.51 CA 9.8 SA i) 32.1 FA 3.93 CA 22.7 SA	a) 10.2% FA 35% CA 102% SA b) 2.9% FA 5.9% CA 45% SA c) 2.5% FA 6.0% CA 27% SA d) 2.9% FA 6.9% CA 33% SA e) 3.4% FA 9.2% CA 36% SA f) 3.1% FA 13% CA 25% SA g) 3.0% FA 12% CA 40% SA h) 3.0% FA 16% CA 20% SA i) 5.1% FA 15% CA 60% SA
Mandak &	Wheat	a) Whole grain bread	- Steryl ferulates (SF)	µg/g dw SF a) 51.2	n/a	µg/g (calculated)	a) 0.01%

Nystrom (2013) ⁷⁷ (Static)		b) Whole grain bread with xylanase c) Whole grain bread with cellulase d) Whole grain bread with xylanase and cellulase e) Baking flour based bread f) Baking flour based bread with xylanase g) Baking flour based bread with cellulase h) Baking flour based bread with xylanase and cellulase		b) 53.0 c) 52.7 d) 52.1 e) 21.7 f) 18.3 g) 19.6 h) 17.0		a) 0.005 b) 0.016 c) 0.016 d) 0.130 e) 0.020 f) 0.004 g) 0.010 h) 0.017	b) 0.03% c) 0.03% d) 0.25% e) 0.09% f) 0.02% g) 0.05% h) 0.10%
Koistinen <i>et al.</i> (2017) ⁷⁶ (Static)	Wheat, rye	a) Bread with native rye bran b) Bread with bioprocessed (enzymatic treatment and fermentation) rye bran	- FA - CA - SA	mg/g a) 1.082 FA 0.037 CA 0.242 SA b) 1.188 FA 0.036 CA 0.258 SA	mg/g a) 0.016 FA 0.001 CA 0.008 SA b) 0.162 FA 0.004 CA 0.029 SA	mg/g absorbed a) 0.549 FA 0.031 CA 0.146 SA b) 1.051 FA 0.034 CA 0.236 SA	a) 51% FA 84% CA 60% SA b) 88% FA 94% CA 91%SA

CA, *p*-coumaric acid; CAF: caffeic acid; FA, ferulic acid; GAE, gallic acid equivalents; PA, phenolic acid; SA, sinapic acid; TPC, total phenolic acid content; dw: dry weight.

‡ as measured in the bread pre-digestion, unless otherwise indicated

* % of bioaccessibility was calculated as the percentage of phenolic compounds in the residue after *in vitro* digestion compared to the initial amount of total PAs/TPC in bread

Table 3. Human studies investigating the bioavailability* and the recovery of bread-derived polyphenols

Reference	Test Samples	Type of study	Subjects	Analysis	Findings
<i>Single-dose dietary intervention</i>					
Bresciani <i>et al.</i> (2016) ⁹⁴	<ul style="list-style-type: none"> - WGB: 94 g of wholegrain bread, 0.926 mg/g total FA; - AB-94: 94 g of a commercial wheat bread enriched in aleurone fraction (6% w/w), 0.458 mg/g total FA; - AB-190: 190 g of a commercial bread enriched in aleurone fraction (6% w/w), ~ 0.458 mg/g total FA. 	Randomized, crossover, single-dose, single-blind, intervention, at least 1-week washout period.	15 healthy subjects, mean age 26 ± 4 y, mean BMI 21 ± 3 kg/m ² .	<ul style="list-style-type: none"> - Plasma ferulic acid- 4'-O-sulfate, dihydroferulic acid-4'-O-sulfate: 0, 0.5, 1, 2, 4, 7 and 24 h; - Urinary ferulic acid-4'-O-sulfate, dihydroferulic acid-4'-O-sulfate, and dihydroferulic acid-O-glucuronide, feruloylglycine, dihydrocaffeic acid sulfate, sinapic acid sulfate, vanillic acid-4-O-sulfate and hydroxybenzoic acid sulfate : 0-3, 3-6, 6-10, 10-14, 14-24, 24-28, 28-34 and 34-48 h. 	<p>Plasma phenolic acid metabolites:</p> <ul style="list-style-type: none"> - Ferulic acid- 4'-O-sulfate C_{max} <ul style="list-style-type: none"> - WGB 84.3 nM; - AB-94 55.5 nM; - AB-190 76.6 nM. - Dihydroferulic acid-4'-O-sulfate C_{max} <ul style="list-style-type: none"> - WGB 9.2 nM; - AB-94 9.5 nM; - AB-190 11.9 nM. <p>- No significantly differences in C_{max} among the tested bread for ferulic acid- 4'-O-sulfate and dihydroferulic acid-4'-O-sulfate.</p> <p>Urine metabolites:</p> <ul style="list-style-type: none"> - Cumulative 48 h excretion <ul style="list-style-type: none"> - Dihydrocaffeic acid sulfate: <ul style="list-style-type: none"> - AB-94: ~2 µmol; - AB-190: ~2 µmol; - WGB: ~0.8 µmol. - Sinapic acid sulfate: <ul style="list-style-type: none"> - AB-94: ~2 µmol; - AB-190: ~1 µmol; - WGB: ~1 µmol. - Significantly higher (<i>p</i> < 0.05) cumulative 48 h excretion of dihydrocaffeic acid sulfate in AB-94 and AB-190 compared to WGB; no statistical differences between AB breads; - Significantly higher (<i>p</i> < 0.05) cumulative excretion of sinapic acid sulfate in AB-190 compared to AB-90 and WGB; no statistical differences between AB-90 and WGB breads. - % Bioavailability: <ul style="list-style-type: none"> - AB-94 +8%; - AB-190: +4%;

					<ul style="list-style-type: none"> - WGB: +4%. - 2-fold higher ($p < 0.05$) bioavailability of the sum of FA in AB-94 compared to WGB and AB-190 WGB. - ~2-fold higher bioavailability of the sum of FA in AB-190 compared to WGB (not significant).
Lappi <i>et al.</i> (2013) ⁹⁵	<ul style="list-style-type: none"> - R bread: 123g commercial wholegrain rye bread (100% rye flour), 0.602 mg/g FA; - WW bread: 109 g white wheat bread, 0.606 mg/g FA; - RB + WW bread: 164 g white wheat bread fortified with native rye bran (35% replacement), 0.713 mg/g FA; - BRB +WW bread: 166 g white wheat bread fortified with bioprocessed rye bran (35% replacement), 0.811 mg/g FA; 	Randomized, cross-over, single-dose, intervention, at least 3-day washout period.	15 healthy subjects, mean age 57 y, mean BMI 26 kg/m ² .	<ul style="list-style-type: none"> - Urinary FA, SA and PA equivalents: the 0–4, 4–12, and 12–24 h. 	<p>FA equivalents bioavailability:</p> <ul style="list-style-type: none"> - BRB+WW: 1%; - RB+WW: 0.4%; - R: 0.4%; - WW: 3.2%. <p>SA equivalent bioavailability:</p> <ul style="list-style-type: none"> - BRB+WW: 0.6%; - RB+WW: 0.4%; - R: 0.3%; - WW: 2.8%. <p>PA equivalent bioavailability:</p> <ul style="list-style-type: none"> - BRB+WW: 0.3%; - RB+WW: 0.3%; - R: 0.3%, 0.07; - WW: 3.8%.
Mateo Anson <i>et al.</i> (2011) ⁴⁹	<ul style="list-style-type: none"> - Control Bread: 300 g whole wheat bread containing native bran, 0.767 mg/g FA, 0.057 mg/g sinapic acid, 0.018 mg/g <i>p</i>-coumaric acid, 0.017 mg/g vanillic acid; - Bioprocessed bread: 300 g bioprocessed bran, 0.733 mg/g FA, 0.057 mg/g, sinapic acid, 0.015 mg/g <i>p</i>-coumaric acid, 0.018 mg/g vanillic acid; 	Randomized, single-blind, single dose, cross-over intervention, at least 1-week washout period.	8 healthy men, range age 21-55 y, range BMI 20-30 kg/m ² .	<ul style="list-style-type: none"> - Plasma ferulic, vanillic and 3,4-dimethoxybenzoic acids relative bioavailability (AUC_{0-t}): 0.25, 0.5, 0.75, 1, 1.25, 1.5, 2, 3, 4, 5, 6, 9, 12 and 24 h. - Urinary FA, SA, CA, VA, and their secondary metabolites: 0 and 24 h. 	<p>Plasma metabolites:</p> <ul style="list-style-type: none"> - FA relative bioavailability (AUC_{0-t}): <ul style="list-style-type: none"> - Control Bread: 240 μmol*min/L; - Bioprocessed Bread: 640 μmol*min/L. - VA relative bioavailability (AUC_{0-t}): <ul style="list-style-type: none"> - Control Bread: 39 μmol*min/L; - Bioprocessed Bread: 70 μmol*min/L. - 3,4-dimethoxybenzoic acid relative bioavailability (AUC_{0-t}): <ul style="list-style-type: none"> - Control Bread: 5.4 μmol*min/L - Bioprocessed Bread: 9.9 μmol*min/L - Significantly higher ($p < 0.05$) relative bioavailability (AUC_{0-t}) of ferulic acid (2.7-fold), vanillic and 3,4-dimethoxybenzoic acid (1.8-fold each) from bioprocessed bread compared to the control bread. <p>Urine metabolites:</p> <ul style="list-style-type: none"> - % Recovery FA: <ul style="list-style-type: none"> - Control Bread: 4%;

					<ul style="list-style-type: none"> - Bioprocessed Bread: 10%. - % Recovery SA: <ul style="list-style-type: none"> - Control Bread: 7%; - Bioprocessed Bread: 15%. - % Recovery CA: <ul style="list-style-type: none"> - Control Bread: 2%; - Bioprocessed Bread: 2%. - % Recovery VA: <ul style="list-style-type: none"> - Control Bread: 104%; - Bioprocessed Bread: 160%. - 2-fold significantly higher ($p < 0.05$) urinary bioavailability of FA, SA, VA, from bioprocessed bread compared to control bread. - No differences in urinary bioavailability of CA from the tested breads.
Chronic dietary intervention					
Harder <i>et al.</i> (2004) ⁹⁶	<ul style="list-style-type: none"> - 250 g control wheat products (Vitacell®), 0 mg FA); - 250 g rye bran enriched products, 0.041 mg/g FA from rye bran - Both the categories included bread, muffin and crisp bread products. 	Randomized, crossover intervention, two 6-week interventions, 4-week washout period.	18 healthy postmenopausal women, mean age 63.3±1.2 y, mean BMI 25.1±0.9 kg/m ²	- Urinary FA equivalents: 0-48 h.	<ul style="list-style-type: none"> - Urinary FA equivalents 24 h-excretion: <ul style="list-style-type: none"> - Baseline: 1.92 mg; - Control wheat: 1.94 mg; - Rye Bran: 4.82 mg; - 1.5-fold higher urinary FA equivalents excretion from rye bran enriched products compared to the baseline (+39.8%, $p = 0.002$) and Vitacell® (+40.2%, $p = 0.001$); - Not significant difference in FA equivalents urinary excretion from Vitacell® products compared to baseline (+1%).
Juntunen <i>et al.</i> (2000) ⁹⁷	<ul style="list-style-type: none"> -Wheat bread consumption (Lignans: 0.109 µg/g); -Rye bread consumption (Lignans, 0.888 µg/g). <p>A minimum of 4-5 slices of bread consumption per day was required, no maximum intake indicated.</p>	Randomized, crossover, 2-week run-in, two 4-week interventions, 4-week wash-out.	43 healthy volunteers, mean age 43±2 y, range BMI 20-32 kg/m ²	<ul style="list-style-type: none"> - Serum ENL concentration: 0 and 4 weeks; - 24 h urinary ENL excretion: 0 and 4 weeks. - 24 h urinary ENL concentration: 0 and 4 weeks. 	<ul style="list-style-type: none"> - Serum ENL concentration <ul style="list-style-type: none"> - Baseline: <ul style="list-style-type: none"> - Men: 28.1 nM; - Women: 39.3 nM. - Wheat bread: <ul style="list-style-type: none"> - Men: 12.5 nM; - Women: 14.8 nM. - Rye bread: <ul style="list-style-type: none"> - Men: 25.6 nM; - Women: 39.7 nM. - Significant higher serum ENL concentrations at the end of rye-brad intervention compared to wheat bread one (+51.2% for men, +62.7% for woman, $p < 0.05$).

					<ul style="list-style-type: none"> - Not significant differences in serum ENL concentration at the end of rye- and wheat-brad compared to baseline. - Urinary ENL 24 h-excretion <ul style="list-style-type: none"> - Wheat bread: <ul style="list-style-type: none"> - Men: 4.0 μmol; - Women: 3.7 μmol. - Rye bread: <ul style="list-style-type: none"> - Men: 6.8 μmol; - Women: 7.8 μmol. - Significantly higher ($p < 0.05$) ENL 24 h-excretion in rye- compared to wheat-bread periods in both men and women. - No correlation between urine ENL and rye bread intake.
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AUC_{0-t} : area under the curve; C_{max} : maximum plasma concentration; CA, *p*-coumaric acid; ENL: enterolactones; FA, ferulic acid; SA, sinapic acid; VA, vanillic acid.

* % of bioavailability was calculated as % ratio between the amount of the compound in the biological fluid on the amount of the ingested compound.

