

1 **The periodic table of fermented foods: limitations and opportunities**

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15 **Abstract**

16 Fermentation is one of the oldest methods of food processing and accounts for a substantial
17 proportion of human foods, including not only staple foods such as bread, cereal porridges or
18 fermented legumes but also fermented vegetables, meats, fish and dairy, alcoholic beverages as
19 well as coffee, cocoa and condiments such as vinegar, soy sauce and fish sauces. Adding the
20 regional varieties to these diverse product categories makes for an almost immeasurable diversity
21 of fermented foods. The periodic table of fermented foods aims to map this diversity on the 118
22 entries of the periodic table of chemical elements. While the table fails to represent the diversity
23 of fermented foods, it represents major fermentation substrates, product categories, fermentation
24 processes and fermentation organisms. This communication addresses limitations of the graphical
25 display on a “periodic table of fermented foods”, but also identifies opportunities that relate to
26 questions that are facilitated by this graphical presentation: on the origin and purpose of food
27 fermentation, which fermented foods represent “indigenous” foods, differences and similarities in
28 the assembly of microbial communities in different fermentations, differences in the global
29 preferences for food fermentation, the link between microbial diversity, fermentation time and
30 product properties, and opportunities of using traditional food fermentations as template for
31 development of new products.

32 **Keywords.** Fermented foods, *Lactobacillus*, *Saccharomyces cerevisiae*, indigenous fermented
33 foods.

34

35 **Key points.**

- 36 • Fermented foods are produced in an almost immeasurable diversity
- 37 • Fermented foods were mapped on a periodic table of fermented foods
- 38 • This table facilitates identification of communalities and differences of products

39

40 **Introduction**

41 Fermented foods have been defined as “foods made through desired microbial growth and
42 enzymatic conversions of food components” (Marco et al. 2021), which emphasizes microbial
43 conversions as the defining characteristic of fermented foods. Fermented foods account for a
44 considerable portion of foods eaten by humans, including not only staple foods such as bread,
45 cereal porridges or beverages and fermented legumes or legume proteins but also fermented meats,
46 fish and dairy, fermented vegetables and alcoholic beverages as well as specialty products and
47 condiments such as vinegar, coffee, cocoa, soy sauce and fish sauces (Steinkraus 1997; Hutkins
48 2019). Fermentation is one of the oldest methods of food processing and has been used since the
49 Neolithic revolution, the transition from hunter-gatherer societies to agricultural societies about
50 14,000 years ago (Hayden et al. 2013; Arranz-Otaegui et al. 2018). Among the fermented foods,
51 the cereal products bread and beer are the oldest fermented food products for which archeological
52 evidence is available (Hayden et al. 2013; Arranz-Otaegui et al. 2018). Just about any agricultural
53 crop or animal product including fruits, cereals, vegetables, milk, fish and meats is fermented at
54 some place in the world, with insect protein as one of few commodities for which traditional
55 fermentation processes have not been described.

56 The microbiology of food fermentation, which initiated the transition from traditional, indigenous
57 knowledge systems to scientific knowledge systems for production of fermented foods, was first
58 described in 1857, when Louis Pasteur attributed alcoholic fermentation to *Saccharomyces*
59 *cerevisiae* (Pasteur, 1857 as reproduced by (Brock 1992). The industrial production of baker’s
60 yeast for bread leavening started in Vienna only 10 years later (Gélinas 2010). Lactic acid bacteria
61 were first isolated by Joseph Lister in the 1870ties (Lister 1877); a comprehensive description of
62 lactic acid bacteria in food fermentations, which remained relevant for much of the 20th century,

63 was published in 1919 (Orla-Jensen 1919). Undefined bacterial starter cultures for baking and
64 dairy fermentations have been available since the late 19th or early 20th century (Brandt 2007);
65 followed by defined strain cultures for dairy, meat, wine, and vegetable fermentations. Food
66 cultures are not only of economic importance but have also been recognized in relation to their
67 beneficial effect on human health (Marco et al. 2017; Wastyk et al. 2021), as microbial cell
68 factories (Sun et al. 2015) and as model systems to study ecology, physiology, and evolution, and
69 domestication of microbes (Wolfe and Dutton 2015; Gallone et al. 2016; Duar et al. 2017b).
70 Evidence for domestication of eukaryotic food fermenting microbes has been provided by large
71 scale comparative genomic analyses of several fermentation organisms including *Saccharomyces*
72 *cerevisiae*, *Aspergillus oryzae* and *Penicillium roqueforti*. In eukaryotes, domestication resulted in
73 distinct phylogenetic clades that are composed exclusively of isolates from food fermentations
74 with a long history of back-slopping. These isolates also exhibit physiological and genetic traits
75 that differentiate the strains from their “wild” ancestors (Gibbons et al. 2012; Gallone et al. 2016;
76 Dumas et al. 2020). Evidence for domestication of prokaryotes, however, remains much less
77 convincing (van de Guchte et al. 2006; Zheng et al. 2015; Kelleher et al. 2017).

78 Despite the economic impact, cultural significance and scientific relevance of fermented foods,
79 only few dedicated textbooks provide an overview on the diversity of fermented foods (Gänzle
80 2019; Hutkins 2019). Since 2014, I have started to map the diversity of fermented foods on the
81 template of the periodic table of chemical elements, a process that was initiated over a Friday
82 afternoon discussion with colleagues that explored whether a “Periodic Table of Fermented Foods”
83 may be a useful tool for teaching of the science of food fermentations at the University of Alberta.
84 An initial version of the periodic table of fermented foods was published in 2015 (Gänzle 2015)
85 but the table continues to be modified with input from collaborators, colleagues and students in

86 the Nutrition and Food Science undergraduate program of the University of Alberta. This
87 communication will briefly present the Periodic Table of Fermented Foods and its outline its
88 limitations. It also aims to outline whether graphical presentation of the diversity of fermented
89 foods to match the periodic table of chemical elements gives rise to relevant scientific questions
90 and hypotheses.

91 **Limitations of the Periodic Table of Fermented Foods.**

92 The periodic table of chemical elements represents a natural law of the periodicity of the properties
93 of the chemical elements (Balarew 2019). In contrast, a natural law of the periodicity of fermented
94 foods does not exist. Moreover, despite all efforts to reduce the author's ignorance on the diversity
95 of fermented foods, to prioritize those fermented foods for which data is available in the scientific
96 literature, and to avoid cultural or geographic bias in the selection of foods that are represented on
97 the periodic table, the presentation remains incomplete and the selection remains to some extent
98 arbitrary. A table with 118 entries cannot represent the diversity of fermented foods. To provide
99 three examples; the periodic table lists 21 cheeses; however, France alone is thought to produce
100 more than 1000 distinct cheese varieties and a comparable number of varieties is produced other
101 countries with a tradition of cheese making (e.g. <http://www.formaggio.it/>). Africa is represented
102 on the table with only 10 entries, however, a continent on which 2000 different languages are
103 spoken can be expected to have as many, or more, different fermented foods. Vegetables that are
104 fermented in a brine with about 2% NaCl are represented by a single entry, sauerkraut, but many
105 different products that employ a comparable fermentation process but use different ingredients,
106 spices or condiments and have different designations are widely consumed in Europe, South Asia
107 and East Asia (Table 1).

108 **Opportunities of organizing the diversity in a Periodic Table of Fermented Foods.**

109 What is the merit, then, of omission of a majority of fermented foods for display of 118 entries in
110 a “periodic table”? First, the process of omission forces to emphasize similarities of different
111 products over differences. For example, fermented vegetables are produced by cutting, brining in
112 salt solutions, followed by fermentation in a closed container at ambient temperature (Ashaolu and
113 Reale 2020). The best known products are sauerkraut, which represents this product category in
114 Figure 1, and kimchi, but numerous other products are produced with similar methods and with
115 comparable fermentation organisms in Europe and Asia (Table 1). Likewise, *mahewu* is produced
116 in Zimbabwe by inoculating a slurry prepared from cooked corn flour with millet malt as a source
117 of amylolytic and proteolytic enzymes, and of fermentation microbiota (Pswarayi and Gänzle
118 2019). Comparable processes and principles are used to prepare cereal beverages in other African
119 countries (Table 2)(Nout 2009; Franz et al. 2014). From a culinary and cultural perspective, these
120 products are very different; from the perspective of the fermentation process or the principles of
121 the assembly of microbial communities, these products share enough similarities to warrant only
122 a single entry that represents all comparable products.

123 A graphical display with 118 entries also allows a quick overview on fermentation substrates,
124 products and fermentation organisms that cannot be provided in a larger figure with several
125 thousand entries to fully map the diversity of fermented foods. Some of these questions – which
126 fermentations include acetic acid bacteria? Which type of products are produced from fish? How
127 is red wine produced from colorless cereal flours? - are readily answered by consulting Figure 1.
128 Other questions that arise from a representative overview rather than a comprehensive display are
129 discussed in more detail below.

130 **Origin and purpose of food fermentation**

131 Publications on fermented foods emphasize the aspect of food preservation and food safety as a
132 motivation for fermenting foods (Nout and Motarjemi 1997; Steinkraus 1997). Preservation is
133 indeed a major aspect in the fermentation of vegetables, dairy products, fish, and fermented meats
134 (groups 12 to 18 in Figure 1). Food fermentations preserve vegetables as a source of vitamins in
135 winter, when fresh vegetables are not available in temperate climates. Fermentation also converts
136 the perishable animal products milk, meat and fish to commodities that can stored and traded over
137 long distances (Kindstedt 2012). Animal agriculture emerged, however, several thousand years
138 after the cultivation and fermentation of the cereal crops (Rowley-Conwy 2011; Arranz-Otaegui
139 et al. 2018) and preservation is thus an unlikely driver for the first food fermentations during the
140 Neolithic revolution.

141 A second major reason for fermentation of fruits, tubers and cereals is the human desire for
142 intoxication, which motivates production of alcoholic beverages in virtually all cultures and on all
143 continents (groups 1 through 4, marihuana edibles and spirits in Figure 1) and has been proposed
144 to be one of the drivers for the first food fermentations in the Natufian (Hayden et al. 2013).
145 Because beer is also a source of energy and nearly isotonic, hydration with low-alcohol beer may
146 be advantageous over water. The notion that medieval city dwellers consumed beer with low
147 ethanol content to avoid contaminated drinking water, however, was identified as a myth
148 (Mortimer 2009; Fusco et al. 2019).

149 A third motivation for food fermentations is the facilitation of milling of cereals and removal of
150 anti-nutritive compounds including phenolic compounds, enzyme inhibitors and phytate, and an
151 improved digestibility of plant crops (Gänzle 2020). Replacing a diverse hunter-gatherer diet with
152 cereal grains, legume seeds, and tubers is a poor proposition unless the palatability and digestibility

153 of these crops and the availability of micro-nutrients is improved by milling, fermentation and
154 heating (Kayodé et al. 2013; Montemurro et al. 2019). In some cases, fermentation is a necessity
155 to remove toxic plant secondary metabolites, e.g. cyanogenic β -glucosides in cassava (group 5),
156 which cause debilitating disease unless they are removed by fermentation or other suitable
157 processes (Kobawila et al. 2005; Nzwalo and Cliff 2011). Steeping of grains also reduces the effort
158 that is needed for wet milling of the grains (Gänzle and Salovaara 2019); an advantage that remains
159 relevant in areas where cereals are processed at the household level. Examples include mawé and
160 ogi produced in Benin (Houghouigan et al. 1993; Greppi et al. 2013) and koko and kenkey
161 produced in Ghana (Halm et al. 1993; Lei and Jakobsen 2004). Extended steeping of cereal grains
162 not only facilitates wet milling but also initiates fermentation, which continues after the milled
163 grains are further processed to porridges or beverages.

164 Last but not least, fermented foods such as miso, soy sauce, vinegar, coffee or vanilla are produced
165 with the purpose to please the palate. While some of the products, e.g. soy sauce analogues or
166 coffee, can be produced with alternative enzymatic or chemical processes that do not involve
167 microbial conversions (Suzuki et al. 2017), fermented products avoid the use of ingredients or
168 additives, and often have superior sensory properties.

169 **Few, if any, of the fermented foods are “ethnic” but almost all represent “indigenous” foods**

170 Figure 1 represents the geographic origin of each product in the upper right of each box, which is
171 colored green if lactic acid bacteria are major members of fermentation microbiota. In recognition
172 to the link of food fermentations to geographic locations and cultures, numerous publications refer
173 to fermented foods outside of Europe and North America as “ethnic fermented foods” (Kwon
174 2015) or “indigenous fermented foods”. The term “ethnic foods” was defined as “an ethnic group’s
175 or a country’s cuisine” (Kwon 2015) or, in a narrower ethnographic meaning, as “food prepared

176 or consumed by members of an ethnic group as a manifestation of its ethnicity” (Anonymous) and
177 thus includes ancestry in the definition of ethnicity. The term “indigenous”, defined by the
178 Merriam-Webster online Dictionary as “produced (...) in a particular region or environment”, is
179 not specific to nations, countries or ethnicity but accommodates fermented foods that are produced
180 by specific to a particular town or other narrowly defined geographic locations that may or may
181 not relate to ethnicity or nationality.

182 A vast majority of fermented foods are specific to, or originate from, specific cultures or regions
183 (Figure 1) and are in some cases a matter of fierce national pride (Jang et al. 2015). This strong
184 link to geography is determined by climate and geography, which determine the availability of
185 fermentation substrates; by the economic constitution of societies, which determines whether
186 fermented foods are produced at the household level, by trades, or in large industrial operations;
187 and by local cultural or religious traditions that define the indigenous knowledge systems on which
188 the fermentation processes are based and the cultural or social context in which fermented foods
189 are consumed (Ströbele 2010). Most fermented foods were produced before scientific knowledge
190 systems were applied for food production. With few exceptions that are discussed below,
191 fermented foods can thus generally be designated as “indigenous foods”.

192 **Community assembly in food fermentations: Differences and similarities between different**
193 **fermentations.**

194 Color coding of the fermented foods informs on the main groups of fermentation organisms;
195 representative microbial species are also indicated. The diversity of fermented foods is matched
196 by the diversity of fermentation organisms. In 2022, an inventory of food cultures with beneficial
197 technological use that has been compiled by the International Dairy Federation included more than
198 226 bacterial and 95 fungal species (Bourdichon et al. 2012; Bourdichon et al. 2022). The

199 characterization of fermentation microbiota by full shotgun metagenomic sequencing (Cao et al.
200 2017) and the description of more than 100 new species of food-fermenting organisms
201 (www.lactobacillus.ualberta.ca/)(Zheng et al. 2020) continues to increase the known diversity of
202 food cultures.

203 Despite this large diversity of fermentation organisms, common patterns for community assembly
204 can be derived from the periodic table of fermented foods. The assembly of communities of
205 organisms is determined by dispersal, selection, speciation and drift (Vellend 2010). Of these four,
206 drift, designating random events, can be ignored if the totality of fermentations rather than an
207 individual fermentation batch is considered. The relevance of dispersal depends on whether the
208 fermentations is controlled by back-slopping (thick box outline and underlined product name in
209 Figure 1)(Li and Gänzle 2020) or relies on the microorganisms that are associated with the raw
210 materials or the processing environment (Miller et al. 2019; Pswarayi and Gänzle 2019). Back-
211 slopping eliminates dispersal limitation and allows recruitment of highly specialized and niche-
212 adapted fermentation organisms (Gänzle and Zheng 2019; Marco et al. 2021). Examples include
213 the host adapted *Streptococcus thermophilus*, *Lactobacillus helveticus* and *L. delbrueckii* in dairy
214 fermentations (Li and Gänzle 2020); the co-existence of *Lactobacillus* and *Limosilactobacillus*
215 species, which is characteristic for the intestinal microbiota of many animals but also observed in
216 back-slopped cereal fermentations (Walter 2008; Duar et al. 2017a; Gänzle and Zheng 2019); and
217 the presence of *Fructilactobacillus sanfranciscensis*, an organism that is likely adapted to insect
218 hosts, in sourdoughs (Gänzle and Zheng 2019). The selective pressure that is exerted by
219 fermentation conditions and raw materials is independent of the geographic location. Each of the
220 three examples indicate above is documented by multiple products from multiple countries
221 representing at least three continents (Figure 1).

222 Spontaneous fermentations that are not controlled by back-slopping or starter cultures also exhibit
223 reliable and globally uniform communities of fermentation microbes that have a stable association
224 with the raw material. This is best exemplified with spontaneous plant fermentations, which are
225 characterized by a consistent succession of fermentation microbiota. Spontaneous plant
226 fermentations are initiated by plant associated *Enterobacteriaceae* including *Cronobacter*,
227 *Kosakonia*, *Klebsiella* and *Citrobacter*, which are among the most abundant representatives of
228 commensal plant microbiota (Schmid et al. 2009; Allahverdi et al. 2016; Pavlova et al. 2017; Taulé
229 et al. 2019). *Enterobacteriaceae* are followed by the more acid tolerant enterococci, lactococci,
230 *Leuconostoc* and *Weissella* species. Eventually the acid tolerant *Lp. plantarum* or pediococci in
231 association with *Lm. fermentum* or *Lv. brevis* prevail (Jung et al. 2012; Wuyts et al. 2018; Pswarayi
232 and Gänzle 2019). This succession of microorganisms is comparable at the family level
233 (*Enterobacteriaceae*) or at the genus level (lactic acid bacteria) for most spontaneous plant
234 fermentations including cereal products or tubers in groups 5 to 7, vegetable fermentations (group
235 12), and coffee and cocoa (Figure 1). Community assembly can be manipulated by addition of salt
236 (e.g. Fu-Tsaii, # 112)(Chao et al. 2009) or by addition of acids to inhibit the initial growth of
237 *Enterobacteriaceae*. Convergence of fermentation communities is also observed for alcoholic
238 fermentations, which all include *Saccharomyces cerevisiae* as a major fermentation organism. All
239 fermentations that include addition of more than 10% NaCl include, irrespective of the substrate,
240 include *Tetragenococcus halophilus* (Figure 1).

241 Speciation or domestication of bacterial species in food fermentation organisms has not been
242 convincingly demonstrated. Although the molecular clock of bacterial evolution is poorly
243 calibrated (Duchêne et al. 2016), the domestication of bacterial organisms with genetic and
244 physiological traits that differentiate fermentation organisms from their “wild” ancestors likely

245 requires more time than elapsed since the onset of back-slopped food fermentations (Duar et al.
246 2017b). Eukaryotes evolve with different mechanisms and at a different pace, though, and
247 domestication of food fermenting yeasts and fungi was demonstrated for *Aspergillus oryzae* from
248 koji fermentation, *S. cerevisiae* from beer and sourdough, and for *Penicillium roqueforti* (Gibbons
249 et al. 2012; Gallone et al. 2016; Dumas et al. 2020; Bigey et al. 2021).

250 In short, the comparison of fermentation microbiota in different fermented foods worldwide
251 demonstrates that, while the fermented products have a strong link to specific regions or countries,
252 the composition of fermentation organisms is globally uniform if comparable substrates and
253 fermentation processes are employed.

254 **North and South, East and West**

255 The periodic table of fermented foods highlights preferences for fermentation substrates and
256 fermentation processes at a global scale (Fig. 1 and Fig. 2). Bread has traditionally been produced
257 in all temperate climates that support cultivation of wheat or rye (Gänzle and Zheng 2019; Arora
258 et al. 2021). In East Asia, steamed bread is preferred (Yan et al. 2019); South Asia, the Middle
259 East and North Africa traditionally produce flat breads; in Europe, bread is baked in loaves.
260 Conversely, fermented cereal foods in Sub-Saharan Africa are consumed predominantly as
261 porridges or non-alcoholic beverages, which are not as common in other parts of the world (Fig.
262 2) (Nout 2009; Franz et al. 2014). The color coding in Figure 2 accounts for the documentation
263 that fermentation cultures that are used in the Americas and Oceania are “immigrants” that were
264 brought by the European that settled on these continents (Salama et al. 1991; Gallone et al. 2016).

265 In Europe and Africa, starch saccharification to produce alcoholic beverages, non-alcoholic
266 beverages or vinegar is achieved by the use of malt; barley malt in Europe or millet and sorghum
267 malts in Africa. In East Asia, starch saccharification is achieved by microbial saccharification

268 cultures. *Koji*, a back-slopped and domesticated cultures of *Aspergillus soyae* or *Aspergillus*
269 *oryzae*, is used in Japan (Gibbons et al. 2012). *Daqu*, a spontaneous fermentation that recruits
270 bacilli, plant-associated *Enterobacteriaceae* and lactic acid bacteria as well as yeasts and moulds
271 to produce amylases and proteases that hydrolyse starch and proteins in a subsequent mash
272 fermentation, is used in China (Fig. 2)(Zheng et al. 2012; Mu et al. 2014). In addition, the
273 traditional use of *Monascus purpureus* to produce red- or yellow-coloured cereal foods is unique
274 to South-East Asia (Lin et al. 2008). Efforts to use the organisms in fermentations in Europe and
275 North America have stalled as the production of red or yellow pigments is invariably associated
276 with the production of the mycotoxin citrinin (Patakova 2013).

277 Milk has traditionally been used for cheese production in Europe, the Mediterranean, the Middle
278 East and the Eurasian Steppes. Communities in Africa and South Asia ferment milk predominantly
279 to yoghurt and comparable set but not strained dairy products (Jans et al. 2017). Conversely,
280 fermentation of legume (soy) protein to diverse products including *tempe*, *natto*, *sufu* or stinky *tofu*
281 is common on East Asia but not in other regions of the world (Fig. 2)(Han et al. 2004; Nout and
282 Kiers 2005; Inatsu et al. 2006). The use of precipitated soy proteins as fermentation substrate also
283 recruits fermentation organisms that are not observed in other parts of the world, e.g. *Rhizopus*
284 *stolonifer* for production of *tempe* (Nout and Kiers 2005) and *Bacillus subtilis*, which is used for
285 fermentation of *natto* (Tsuji et al. 2015).

286 Fish fermentations have traditionally been used in Scandinavian countries and in South East Asia.
287 Examples for Scandinavian fermentations include *harkarl*, fermented shark produced in Iceland,
288 and *surströmming* in Sweden (Skåra et al. 2015). East Asia produces fermented fish sauces, where
289 the composition of fermentation microbiota is controlled by addition of more than 10% NaCl, and
290 fermented sour fish where the composition of fermentation microbiota is controlled by addition of

291 carbohydrates including starch (rice) or sugars and / or addition of salt (Paludan-Müller et al.
292 2002). The author is unaware of comparable fermented products in other regions of the world
293 where seafood is available. The production of *garum*, however, a fish sauce that was produced in
294 ancient Rome but came out of fashion after the fall of the Roman Empire (Corcoran 1963),
295 indicates that this is a question of preference.

296 **Microbial diversity and product properties**

297 The arrangement of selected fermented foods in the periodic table of fermented foods roughly
298 matches the flavor intensity within each groups, with the blandest examples at the left and top and
299 the product with the most intense flavor at the right and bottom (Figure 1). This arrangement
300 indicates that long fermentation times and / or a diverse fermentation microbiota results in a more
301 intense flavor. The case can be convincingly made with two somewhat exotic examples,
302 *surströmming*, a fermented fish product from Sweden, and kopi luwak (civet coffee). The
303 fermented fish product *surströmming* is produced with diverse fermentation organisms that
304 represent several bacterial phyla including *Firmicutes*, *Bacteroidetes*, *gamma-Proteobacteria* and
305 *Actinobacteria*. The resulting product smells somewhat intense (Belleggia et al. 2020). Kopi
306 Luwak is produced by feeding civet cats with coffee fruits and collecting the excreted beans after
307 fermentation by the intestinal microbiota of civet cats. Intestinal microbiota of the civet cat are
308 dominated by acetic acid bacteria, lactic acid bacteria and *Enterobacteriaceae* (Watanabe et al.
309 2020), which overlaps with those organisms dominating wet fermentation process of coffee beans
310 (de Melo Pereira et al. 2017) but includes additional organisms as well as digestive enzymes of
311 the civet cat.

312 The case can also be made with the more commonly consumed fermented products bread and
313 cheese. Long-term cheese ripening recruits non-starter lactic acid bacteria that contribute to flavor

314 formation in addition to the starter cultures (Lo et al. 2018). During cheese ripening, casein is
315 hydrolysed to taste-active peptides and amino acids, in particular glutamate, which can accumulate
316 to levels exceeding the taste threshold more than 500-fold (Toelstede and Hofmann 2008;
317 Hillmann et al. 2016). Straight-dough bread is fermented only with baker's yeast while sourdough
318 baking includes a contribution of lactic acid bacteria and sourdough yeast, predominantly
319 *Kazachstania humilis*, to biochemical conversions during bread-making, and generally involves
320 extended fermentation times (Gänzle 2014; Gänzle and Zheng 2019; Arora et al. 2021). In
321 comparison to straight dough bread, sourdough bread is characterized by a greater diversity and
322 higher concentration of taste-active compounds and odour volatiles (Hansen and Schieberle 2005;
323 Zhao et al. 2015). A last example relates to the comparison of two distilled grain liquors, whisky
324 and *baijouw*. Whisky is fermented with *S. cerevisiae* with a variable contribution of lactobacilli (van
325 Beek and Priest 2000); odorants are additionally derived from the malt, the peat smoke used for
326 drying of the malt and the casks used for maturation (Jeleń et al. 2019). The fermentation process
327 for grain liquors in China includes contributions from diverse microbes including yeasts, fungi,
328 bacilli, *gamma-Proteobacteria* and *beta-Proteobacteria*, and *Firmicutes* which include but are not
329 limited to *Lactobacillales* (lactic acid bacteria)(Zheng and Han 2016). Again, the higher diversity
330 of microbes that is recruited for *baijouw* fermentation results in a higher diversity and intensity of
331 flavor volatiles (Jeleń et al. 2019; Chen et al. 2021).

332 **Tradition and innovation**

333 The industrialization of food production also resulted in the industrialization of the production of
334 fermented foods. This process generally involved scaling of traditional fermentation processes and
335 transition from traditional, indigenous knowledge systems to scientific knowledge systems; in
336 short, moving from “art to science”. Currently, food fermentations extend to products for which

337 no traditional template exists. In these cases, the periodic table of fermented foods can guide the
338 development of fermentation processes in the absence of a traditional template. Examples include
339 the fermentation of insects (Kewuyemi et al. 2020) or the production of plant cheeses (Jeewanthi
340 and Paik 2018), where information on fermentation of other protein foods (soy, groups 10 and 11,
341 dairy, groups 13 – 16 and fish and meat, groups 17 and 18) may provide useful information on the
342 use of fermentation organisms and enzymes to improve product quality. Moreover, fermentation
343 of vegetables has re-emerged as a method to provide high-quality food not only at the household
344 level but also by chefs and small start-up companies. The corresponding products are not limited
345 to those for which traditional templates exist (Redzepi and Zilber 2018).

346 Current commercial relevance for novel, non-traditional fermented foods relate for example to
347 alcohol-free fermented cereal beverages and gluten free bread. Fermentation of malt with lactic
348 acid bacteria or acetic acid bacteria allows adjusting the level of sweetness, acidity and carbonation
349 or “fizz” to levels that meets consumer’s expectations (Bronnmann and Hoffmann 2017). Non-
350 alcoholic fermented cereal beverages are widely consumed in Africa and, to a lesser extend, in
351 East Europe but not in Central or Western Europe (Taylor 2016). Element # 102, Bionade, provides
352 an example of a non-traditional fermented “designer” food. Likewise, the development of gluten-
353 free bread in the last two decades necessitated building on information related to traditional
354 fermentation of sorghum, millet or corn fermentations to develop a fermented food for which no
355 traditional template is available (Gallagher et al. 2004).

356 **Concluding remarks.**

357 Mapping the diversity of fermented foods on a simple graphical display is impossible in a world
358 that is inhabited by 8 billion humans where many products are fermented at the household or
359 regional level. The effort to produce such a simple graphical display nevertheless has merit as it

360 necessitates acknowledgement of the – largely unknown - diversity but also allows to derive
361 common patterns in the fermentation of products that, at first sight, appear to be very different.
362 Many of the fermented foods contain live fermentation microbes at the time of consumption
363 (groups 6, 7, 12 to 18, and several elements in the group “tea, coffee, chocolate, and various
364 beverages). Live microbes that are present in fermented foods are increasingly recognized as
365 contributors to human health even if no strain specific health claims were established (Marco et
366 al. 2021; Wastyk et al. 2021). The periodic table of fermented foods also provides an indication of
367 the many fermented foods that are likely to please one’s palate but remain to be sampled. Last but
368 not least, many fermented foods negate conventional wisdom and are tasty *and* healthy.

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732

733 **Figure legends**

734 **Figure 1.** Periodic element of the fermented foods. The figure aims to provide an overview of the
735 diversity of fermented foods that are produced globally. Indicated are the origin of the food, main
736 ingredients and typical fermentation organisms, pH, water activity and fermentation / ripening
737 time, and major microbial metabolites that relate to product quality. The contribution of major
738 groups of fermentation organisms is color coded as indicated in the legend; back-slopped
739 fermentations are indicated by a thick outline and by an underlined product name. The figure is
740 not provided in this pre-print; the reader is referred to the updated, high resolution file for printing
741 in A0 format that is available on the author's personal website and on:

742 <https://drive.google.com/open?id=10cH7XaNqGQAKdL3VHhd1tzWXlGaTdfkV&authuser=mg>
743 [aenzle%40ualberta.ca&usp=drive_fs](https://drive.google.com/open?id=10cH7XaNqGQAKdL3VHhd1tzWXlGaTdfkV&authuser=mg)

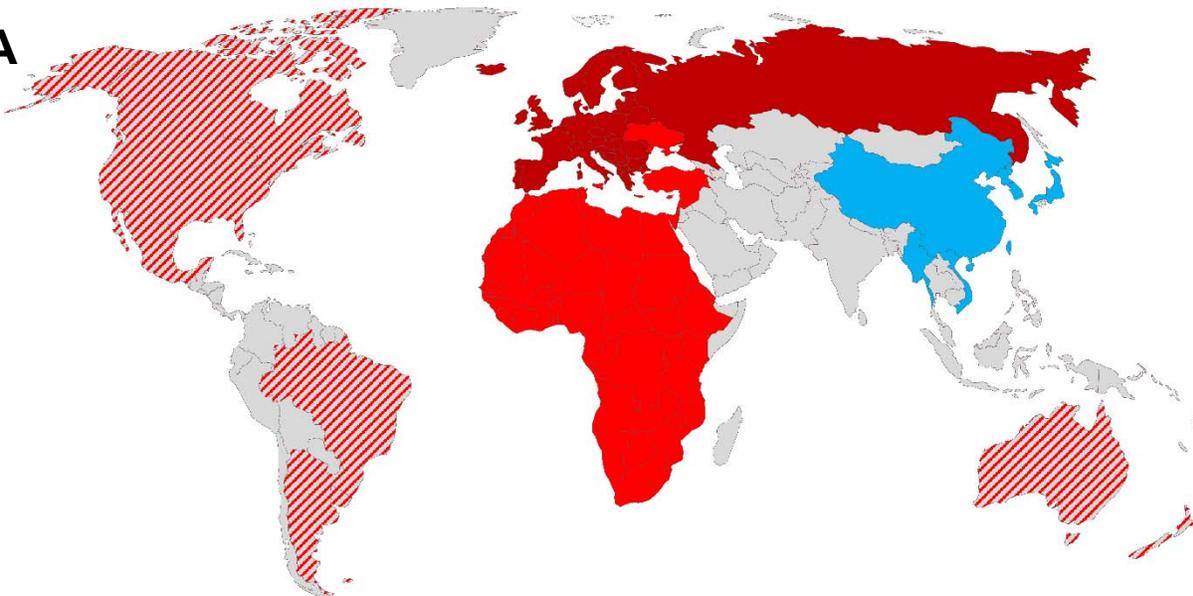
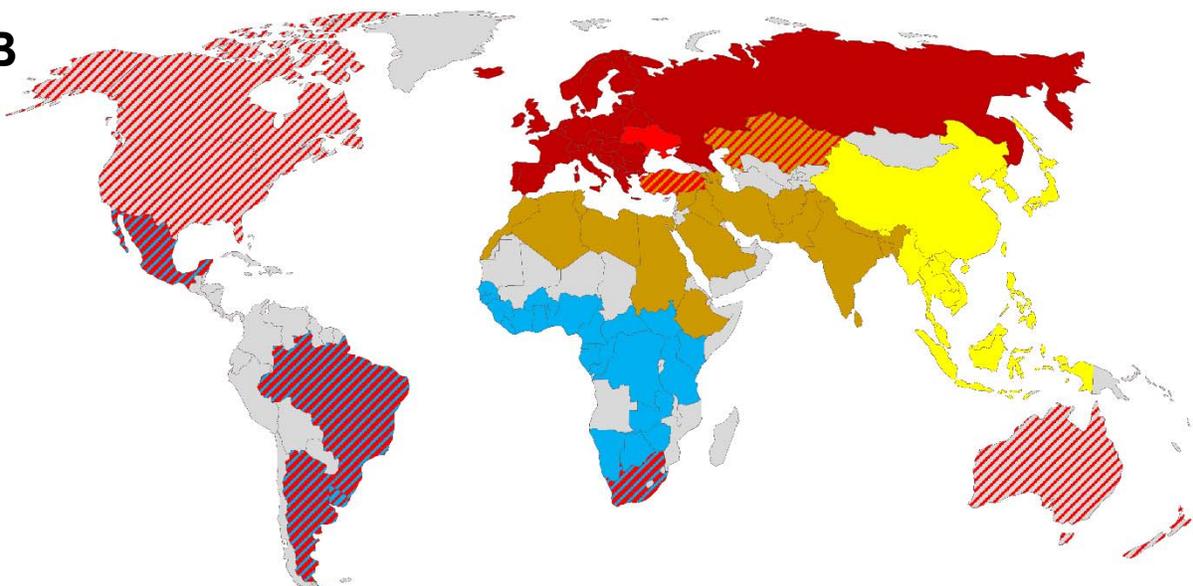
744 **Figure 2.** Global preferences for traditional consumption of different types of fermented foods.
745 Preferences are color coded based national borders, ignoring regional preferences. **Panel A.** Use
746 of malt (red) or microbial enzymes (blue, *koji*, *daqu*) for production of alcoholic beverages or
747 vinegar from grains. **Panel B.** Consumption of bread as steamed bread (yellow), flat bread
748 (orange), in loaves (red) or consumption of cereal porridges or beverages (blue). Countries are
749 hatched if they are at the intersection of two cultures (e.g. Turkey) or if the preferences of
750 indigenous and immigrant (fermentation) cultures differ (e.g. Brazil, North America). **Panel C.**
751 Production of isolated, fermented products from soy (blue) or of cheese (red). Countries that
752 produce little, if any, cheese (most African countries), or produce acid-coagulated cheese without
753 fermentation (South Asia) are coloured in orange. Countries are hatched if preferences of
754 indigenous and immigrant (fermentation) cultures differ. Gray, not relevant or insufficient
755 information. Maps were drawn with a template file from www.freeworldmaps.net.

Table 1. Examples for vegetables that are produced by cutting, brining, and fermentation for 2 – 4 weeks at ambient temperature, matching element #30, sauerkraut.

Name	Origin	Main ingredients	References
Sauerkraut / Choucrute	Germany, France, U.K., U.S.A and Canada	cabbage (juniper berries)	(Hutkins 2019)
Tursu	Turkey	cabbage, cucumber, carrot, beet,	(Çetin 2013)
Gundruk	India	Cabbage, mustard leaves, raddish and / or cauliflower	(Tamang and Tamang 2010)
Sinki (raddish)	India, Nepal	radish	(Tamang et al. 2005)
Pao Cai	China	cabbage (red pepper , scallion , ginger, garlic, anchovy)	(Liu et al. 2019b)
Suan-Tsai, Suan Cai	China, Taiwan	Napa cabbage (North China) or green mustard (South and West China, Taiwan)	(Chao et al. 2009; Liu et al. 2019a)
Jianshui	China	Cabbage or celery	(Jun et al. 2018)
Kimchi	Korea	cabbage (red and black pepper, onions, mustard)	(Patra et al. 2016)
Dua muoi or dha muoi	Vietnam	Mustard or beet	(Nguyen et al. 2013)
Ca muoi	Vietnam	Eggplant	(Nguyen et al. 2013)

Table 2. Examples for non-alcoholic fermented cereal beverages that are produced by fermentation of a cooked cereal slurry with addition of adjuncts as source of enzymes and fermentation microbiota, matching element # 42, mahewu.

Name	Origin	Main ingredients	References
Mahewu, Amahewu, Emahewu, or Mageu	Zimbabwe South Africa, Swaziland	cooked maize with millet or sorghum malt	(Pswarayi and Gänzle 2019)
Togba	Zimbabwe	cooked maize	(Gadaga et al. 1999)
Tobwa	Tanzania	cooked maize with millet or sorghum malt	(Mugula et al. 2003)
Munkoyo (Chimbwantu)	Zambia, Democratic Republic of Congo	cooked maize with millet malt, <i>Rynchosia</i> root, cowpea root or sweet potato peels	(Phiri et al. 2019)
Ekitiribita	Uganda	cooked millet with sorghum malt	(Mukisa et al. 2010; Mukisa et al. 2012)

A**B****C**