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Additional Information

1 **Fat Digestibility in Meat Products: Influence of Food Structure and**
2 **Gastrointestinal Conditions**

3

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27 **Fat Digestibility in Meat Products: Influence of Food Structure and** 28 **Gastrointestinal Conditions**

29 Digestibility of macronutrients depends on the food matrix structure as well as on
30 gastrointestinal conditions, especially in patients with exocrine pancreatic
31 insufficiency. In this situation, an oral enzyme supplementation that promotes
32 nutrient hydrolysis is needed. In this context, in the present study, a static in vitro
33 digestion model was used to assess the lipid digestibility of different meat
34 products (processed and fresh), different intestinal conditions of pH (6 or 7), bile
35 concentration (1 or 10 mM), and doses of the enzyme supplement (1000 to 4000
36 lipase units/ g fat). Results showed that processed (unstructured) meats had better
37 matrix degradation during digestion and reached higher values of lipolysis
38 extents (total free fatty acids/ g fat) than the natural meat matrices with a
39 statistically significant association ($p<0.001$). Regarding the intestinal medium,
40 pH of 7 and bile concentration of 10 mM contribute to higher matrix degradation,
41 and thus, to a higher lipolysis ($p<0.001$).

42
43 Keywords: in vitro digestion, lipolysis, nutrition, pancreatic insufficiency, food
44 matrix

46 **Introduction**

47 Nowadays meat products are overconsumed in the western societies, both in young and
48 adult populations, and the consumption exceeds the recommended daily intake
49 (Huybrechts et al, 2010, Moreira et al, 2010, Pérez-Rodrigo et al. 2015, Roccaldo et al.
50 2014). Processed meats are the preferred choice over the fresh varieties because
51 processed meat products are cheaper and easier to prepare. Moreover, the fast-food type
52 restaurants, which offer mainly the varieties of meat that are already prepared and
53 processed, are a frequent meal choice (Bhutani, Schoeller et al. 2018, Close et al. 2016,
54 French et al. 2001).

55 Meat is an important source of protein and vitamins and its composition also
56 contains fat and water (Pereira and Vicente 2013). Different meats and cuts are
57 composed of distinctive types of muscle fibres and they contain varying amounts of
58 connective tissues. This implies that the quality and structure of them are very diverse
59 (Greenfield et al. 2009). In addition, the processing method of meat, ~~as well as~~

60 including the cooking technique, may also affect the final structure resulting in a wide
61 variety of food matrices in this food group (Bax et al.2012, Luo et al. 2018).
62 Concerning the fat fraction and quality of the meat, they depend on the area of the meat
63 portion as well as the origin of the animal.

64 While several studies assessing proteolysis in meat products have been
65 published in the recent years, scarce literature is available regarding the study of
66 lipolysis. Given the direct effect of dietary fat intake on the development of diet-based
67 diseases (de Souza et al. 2015), it is important to unveil the digestion fate of this
68 nutrient when it comes from a food choice as popular as meat.

69 Processed meats normally have a higher fat content than non-processed
70 varieties, as they are made of the animal parts that are less consumed, have less sensory
71 quality and a lower consumer acceptance. However, apart from the fat content, the
72 structure also could influence digestibility of this nutrient. Concretely, in natural
73 structure of meat, fat molecules are embedded in muscle fibres while processed meats
74 have an unstructured mixture of fat, protein and water, and in some cases, carbohydrates
75 are also added (Guo et al. 2017). Therefore, in order to address the study of lipid
76 digestibility in meat, the influence of the meat structure should be taken into account.

77 Lipid digestion is a complex process, which takes place in the duodenal region
78 of the small intestine, which implies a previous micellization in order to provide the
79 enzyme access allowing lipid hydrolysis. During this process, the intestinal pH and the
80 presence of bile salts are crucial to improve the enzymatic hydrolysis (Ryan et al. 2008,
81 Whitcomb et al. 2010). The intestinal pH is directly related to the enzyme activity and
82 efficiency. Moreover, the bile salts act as emulsifying agents leading to the
83 micellization of fat and providing a larger surface area on the fat globule for the enzyme
84 to adsorb. However, the alteration of these factors could compromise the lipid
85 digestibility in some individuals who suffer from exocrine pancreatic insufficiency
86 (EPI), mainly present in patients with Cystic Fibrosis. The obstruction of the pancreatic
87 duct in EPI produces an insufficient secretion of pancreatic juice, which contains the
88 digestive enzymes. Moreover, the secretion of sodium bicarbonate (NaHCO_3) is also
89 reduced, causing a drop in the intestinal pH and the consequent reduction of the
90 pancreatic enzymes activity. This situation leads to the mal-digestion and mal-
91 absorption of nutrients (Layer and Keller 2003, Naikwade et al. 2009). The treatment
92 for EPI consists in oral administration of pancreatic enzyme supplements, containing
93 proteases and mainly lipase (Armand et al. 2011). The current guidelines for nutritional

94 management in Cystic Fibrosis recommend a dose of the enzymatic supplement of 2000
95 – 4000 LU/ g fat, with a very low degree of evidence (Turck et al. 2016). The
96 modulation of both intestinal pH and bile salts concentration in the intestinal digestion
97 medium are thus of upmost relevance in the process of lipolysis.

98 In vitro studies are a useful tool that allows studying the structural changes, the
99 digestibility and release of food components under simulated gastrointestinal
100 conditions. In this context, the goal of the present study was to evaluate the influence of
101 gastrointestinal conditions associated with EPI (intestinal pH, bile salts concentration
102 and the dose of the enzyme supplement) as well as the effect of the food matrix on lipid
103 digestibility in different meat products.

104

105 **Materials and methods**

106 *Materials*

107 The different meat products (hamburger (just the meat patty, 100% minced pork meat),
108 sausage, luncheon ham, cooked ham, cured ham, pate, chicken drum, pork loin and beef
109 steak) were obtained from a local supermarket in Valencia (Spain).

110 For the preparation of the simulated digestive fluids (**Table 1**) the following
111 chemicals were needed: human α – amylase (1000 – 3000 U/ mg protein) (CAS: 9001-
112 19-8), pepsin from porcine gastric mucosa (≥ 2500 U / g protein) (CAS: 9001-75-6),
113 bovine bile extract (CAS: 8008-63-7), KCl, KH_2PO_4 , NaHCO_3 , NaCl, $\text{MgCl}_2 \cdot (\text{H}_2\text{O})_6$,
114 $(\text{NH}_4)_2\text{CO}_3$ and CaCl_2 , all of them from Sigma-Aldrich Chemical Company (St Louis,
115 MO, USA). NaOH (1 N) and HCl (1 N), were all acquired from AppliChem Panreac.
116 Pancreatic enzymes supplements (Kreon 10,000 lipase units (LU)) were used to
117 simulate in vitro digestion of an individual with EPI. Each capsule contains 150 mg of
118 gastro-resistant microspheres containing porcine pancreatic enzyme equivalent to
119 10,000 lipase U., 8,000 amylase U., and 600 protease U. The specific lipase activity of
120 the Kreon was usually measured before the experiments (Carriere et al. 2000) and the
121 amount of supplement added to the gastric stage was adjusted always to have the
122 corresponding LU/g fat according to the experimental design.

123 For the analytical determinations, Triton-X 100%, as well as the analytical
124 standard of palmitic acid, were acquired from Sigma-Aldrich. Ethanol (96 % v/v for
125 analysis) was acquired from AppliChem Panreac.

126

127 *Sample preparation*

128 Prior to the digestion process, hamburger, chicken drum, pork loin and beef steak were
129 placed in a microwavable plate and cooked in a household microwave oven (model
130 GW72N, Samsung) for 4 min at 600 W, 2450 MHz. For the rest of the matrices
131 (luncheon ham, pate, cured ham, cooked ham and sausage) a thermic treatment was not
132 necessary because they were ready to eat. After their preparation, all samples were in
133 vitro digested by using a static system.

134

135 *In vitro digestion process*

136 Meat samples were placed into falcon tubes (50 ml); the amount of samples to be
137 digested was weighted in order to have 0.35 g of fat in each tube. Fat, water and protein
138 content in all meat products was determined before digestion by the official methods
139 (AOAO, 2000).

140 The digestion proceeding used was based on the standardized static in vitro
141 digestion method for food published by Minekus et al. (2014) with some modifications
142 in order to allow for analysing EPI conditions. **Table 1** illustrates the amounts and
143 composition of the fluids required in each of the stages of the digestion process
144 (Minekus et al. 2014). The digestion fluids were prepared daily from stock solutions,
145 salivary (SSS), gastric (SGS) and intestinal (SIS) prepared according to Minekus et al.
146 (2014). The enzymatic activity was tested before each simulation following the protocol
147 proposed by Carriere et al., (2000). The in vitro digestion process was performed as
148 follows:

149 Oral stage: Simulated salivary fluid (5 ml) (SSF; pH 8) at 37°C, was added to
150 the meat sample in a 1:1 (w/v) ratio and properly homogenized with a kitchen blender
151 for 3 minutes (Vario Mixer, Ufesa 600 W). Salivary amylase was added into SSF to
152 reach a concentration in the saliva mixture of 75 U/ml.

153 Gastric stage: After the oral stage, simulated gastric fluid (SGF; pH 3) was
154 added to each tube containing the oral bolus (1:1 v/v). Pepsin was added into the SGF to
155 reach a concentration in the gastric mixture of 2000 U/ml. The pH of the mixtures was
156 adjusted with HCl (1 N) to $\text{pH } 2.8 \pm 0.1$ and samples were flipped from top to bottom at
157 55 rpm for 2 hours using an Intell-Mixer RM-2 (Elmi Ltd, Riga, LV-1006, Latvia) and
158 incubated at 37 °C in a chamber Selecta (JP Selecta SA, Barcelona). These mixing
159 conditions provided constant mechanical energy to induce the breakdown of the food
160 matrix during digestion. The pancreatic supplement was added in the gastric stage in
161 order to simulate swallowing the pill in case of EPI situations.

162 Intestinal stage: Following the gastric stage, simulated intestinal fluid (SIF; pH
163 7) was added in a 1:1 (v/v) proportion to each tube containing the gastric chime. The pH
164 of the mixtures was adjusted to pH 6.0 ± 0.1 or 7.0 ± 0.1 , depending on the conditions
165 to be tested, with NaOH (1 N). Samples were then being agitated from top to bottom at
166 55 rpm for another 2 hours at 37 °C. pH was monitored during the digestion process and
167 readjusted if necessary to keep it constant (González-Bacerio et al. 2010, Prazeres et al.
168 1994).

169

170 ***Experimental design***

171 The experimental design for each type of product (hamburger, sausage, luncheon ham,
172 cooked ham, cured ham, pate, chicken drum, pork loin and beef steak) consisted of two
173 main sets of experiments. In the first, intestinal conditions were fixed at pH 6 and bile
174 salts concentration 1 mM, and different pancreatin supplement doses (0, 1000, 2000,
175 3000 and 4000 LU/g of lipid) were tested, in order to assess the influence of enzyme
176 concentration. In the second, the dose of enzymes was fixed at 2000 LU/ g of lipid, and
177 the study variables were different combinations of intestinal pH and bile concentration:
178 pH6 – 1 mM, pH 6 – 10 mM, pH 7 – 1 mM and pH 7 – 10 mM, in order to analyse the
179 impact of different intestinal scenarios on lipolysis, and matrix degradation. Of note, the
180 combination pH 6 – 1 mM would represent the most unfavourable condition in the
181 gastrointestinal tract in EPI (Clarke 2001, Gelfond et al. 2013, Harries et al. 1979,
182 Rovner et al. 2013, Vu et al. 2000), and the pH 7 – 10 mM would approach the standard
183 duodenal conditions of a healthy adult. All the experiments were conducted in triplicate.

184

185 ***Analytical determinations***

186 ***Matrix Degradation Index (MDI (%))***

187 Matrix Degradation Index (MDI) was determined in all samples after in vitro digestion.
188 This parameter represents the proportion of solids that were finely dispersed in the final
189 product of digestion. The total content of a digestion tube was centrifuged (4000 x g for
190 20 minutes, 4 °C) and filtered by a metallic sieve (1.6 mm x 1.6 mm mesh) to separate
191 out large particles. The drained liquid was collected and used to determine the free fatty
192 acids. The solid particles were rinsed twice with 5 ml of appropriate juice to remove any
193 digested material. Blotting paper was placed around the metallic sieve for 10 minutes to
194 drain residual digestion juice. The solid meat particles were then transferred to an
195 aluminum dish and immediately weighed. The aluminum dish was put in a forced air

196 oven at 60 °C for 48 hours and weighed again to determine the mass of large solids. The
197 MDI, corresponding to the proportion of solids passing the metallic sieve, was
198 calculated according to Lamothe et al. (2012, 2014).

199 *Lipolysis extent (%)*

200 Drained juice from digested samples was diluted 100-fold with a solution made of 5.6%
201 Triton X-100 and 6% ethanol in water (Lamothe et al. 2012) . This solution was used to
202 solubilize the free fatty acids (FFA) and stop the lipase activity. Fatty acids release
203 during digestion was measured on the diluted samples using a free fatty acid
204 colorimetric assay kit (Roche Diagnostics, Indianapolis, IN, USA) and a
205 spectrophotometer (UV/vis, *Beckman Coulter*) (Lamothe et al. 2014). Palmitic acid
206 standard was used for quantitative determination of FFA. Total FFA was expressed as
207 mg of fatty acids released after a complete digestion per gram of initial fat present in
208 each meat product.

209

210 ***Statistical analysis***

211 The variables included for the statistical analysis were the nutrient composition of
212 foods: water, protein and fat, starch; and the food matrix structure: natural food matrix
213 (protein fibres) or processed matrix (unstructured); the pancreatic enzyme supplement
214 concentration (enzyme dose); the intestinal conditions: pH and bile salts concentration;
215 and the matrix degradation index (MDI). The response variables were lipolysis extent
216 (total FFA) and the MDI.

217 Data were summarised using mean, standard deviation, median and 1st and 3rd
218 quartile in the case of continuous variables and with absolute and relative frequencies in
219 the case of categorical variables.

220 Linear mixed regression models were performed to assess the effect of the food
221 matrix structure and other factors such as matrix degradation index were included as
222 covariates. Additionally, because observations of the same food are more likely to have
223 similar lipolysis extent due to their nutritional characteristics, the linear regression
224 models were extended with the "Food" variable as random effect with random intercept
225 to correct for the non-independence of the data.

226 All the analyses were performed by software R (version 3.4.2) using packages
227 betareg (version 3.1-0), lme4 (version 1.1-14) and NMF (version 0.20.6). A p-value
228 lower than 0.05 was considered statistically significant.

229

230 **3. Results and discussion**

231 *Effect of the food structure on matrix degradation and fat digestibility*

232 The degradation of the food matrix is the process by which the 3D structural
233 conformation of a food is disrupted into smaller parts allowing for the release of the
234 structural components, i.e. the nutrients (**Table 2**). It is facilitated by several
235 mechanisms: mainly the mechanical forces produced along the gastrointestinal tract
236 (chewing in the mouth, stomach walls agitation and small intestine peristaltic
237 movements) and the enzymatic activity that contributes to the breakdown of nutrients
238 conforming the matrix structure. Thus, when no enzymes were used in the in vitro
239 digestion, significantly lower matrix degradation indexes were obtained (**Figure 1**) as
240 compared to digestions conducted with pancreatic enzyme supplements. Generally, the
241 enzymatic supplements increase of 1000 LU/ g of fat led to a minor increase in the
242 MDI. In the case of the natural matrices (protein fibres) maximum MDI were between
243 50 and 75%, while in the unstructured matrices higher degradation extents were reached
244 (>75%).

245 In terms of fat digestibility (total release of FFA), a similar tendency as in the
246 MDI was shown for all the assessed meat products (**Figure 2**). In those matrices which
247 preserve the protein fibres structure, the total FFA increased with the concentration of
248 the enzymatic supplement up to 3000 LU/g fat, and it slightly decreased at 4000 LU/g
249 fat, probably due to inactivation by aggregation. It has been described that when a high
250 concentration of enzymes is not active in the reaction medium (e.g. because there is no
251 substrate available) aggregation and inactivation occur. Furthermore, the release of fat
252 particles from the protein fibres complex is a progressive and slow process, so this may
253 lead to a low concentration of substrate at the beginning of the digestion. This may be
254 the reason of the inactivation phenomenon (López-Gallego et al. 2005). In contrast, in
255 unstructured matrices, the release of FFA was directly proportional to the concentration
256 of enzymes. Pate and luncheon meat reached a total FFA released close to 600 mg/ g fat
257 at enzyme dose of 4000 LU/ g fat. On the other hand, hamburger and sausage reached a
258 lipolysis extent of 500 and 300 mg FFA/ g fat, respectively. In any case, the highest
259 value of lipolysis was reached at enzyme dose of 3000 LU/ g fat.

260 The fact that MDI and total FFA released followed the same tendency was,
261 indeed, statistically explained, finding significant associations between the type of
262 matrix and the MDI, and between the MDI and the fat digestibility (**Figure 3**). The
263 unstructured matrices allowed for higher MDI than the natural protein fibres, up to

264 30.6% more ($p = 0.013$, 95% CI [8.148, 30.612]); and the higher MDI were associated
265 with higher total FFA released, every 1% increase produced between 6.1 and 9.6 mg of
266 FFA more ($p < 0.001$, 95% CI [6.128, 9.606]).

267 These relationships are backed up by previous literature providing a physical
268 explanation, as previously compiled by Guo et al. (2017). When fat molecules are
269 trapped in a solid food matrix, the structure of the surrounding food matrix is the
270 dominant factor controlling digestion. In order to make lipids available to the enzymes,
271 the degradation of the protein food matrix has to occur. In such systems, lipolysis is
272 thus conditioned by the rate and extent of proteolysis: as the protein structure is broken
273 down, the lipids are released from the matrix, and lipases can start hydrolysis
274 (Dickinson 2012). An example of this type of system was described by Dickinson
275 (2012), in which protein gels containing lipids were formulated with different amounts
276 of protein and forming different microstructures (hard and soft gels). The soft gel
277 presented an in vitro digestion behaviour similar to a liquid whey protein emulsion, but
278 the hard gel showed slower lipolysis rates and extents, as the gel structure was degraded
279 along digestion and allowed for fat droplets release from the matrix (Guo et al. 2017,
280 Guo et al. 2016).

281 Another example supporting our findings is the study of fat digestibility in
282 chewed almonds versus their isolated fat molecules, which lipolysis extents were 22 and
283 69% after one hour of in vitro digestion (Grundy et al. 2017). The study pointed at the
284 fact that cells remained largely intact after mastication and concluded that lipid
285 bioaccessibility in almonds was dependent on the structure and the cell walls
286 surrounding the oil droplets.

287 Finally, the role of the food structure in protein-lipid matrices has been also
288 assessed in two types of cheese (Fang et al. 2016). Cheddar cheese had larger fat
289 globules that made the structure less hard and more easily degraded. On the other hand,
290 the mozzarella had a denser fibrous protein matrix. In terms of digestibility, it was
291 higher in cheddar as fat globules were rapidly released and accessible to the enzymes,
292 while in mozzarella the fibrous structure prevented fat release and thus led to a lower
293 lipolysis extent.

294 In the present study, natural protein fibres matrix represents a robust structure in
295 which lipids are contained, and the unstructured matrices suppose a softer system in
296 which lipid and protein are not bonded or linked to each other. This structural fact

297 explains why the unstructured matrices had more MDI and consecutively a higher
298 amount of FFA release.

299 Apart from the food structure, the conditions of the digestive fluid and the
300 intestinal environment may also affect the behaviour of food digestion. These are
301 explored and explained in the coming section.

302

303 *Effect of the intestinal conditions on matrix degradation and fat digestibility*

304 In all the assessed intestinal conditions, the unstructured matrix meat products had
305 higher MDI than the structured, values ranging from 69 to 78.6% and 42 to 70%,
306 respectively (**Table 3**). The intestinal pH had a significant effect on the matrix
307 degradation index, as in all the assessed meat products it allowed for an increase
308 between 5 and 15% ($p < 0.01$). However, when comparing the effect of the bile salts
309 concentration at intestinal pH 6 or at pH 7, this showed a non-significant increase in the
310 matrix degradation index. Pate was an exception, in which MDI increased 7% at the 10
311 mM concentration, both in pH 6 and 7. In contrast, the role of bile was not significant in
312 proteolysis, but it was important on lipolysis due to the bile emulsify the lipids allowing
313 a higher surface area available for the lipases action.

314 Overall, in terms of fat digestibility, the release of total FFA was significantly
315 higher at pH 7 ($p < 0.001$, 95% CI [53.5, 168.9]) and at bile salts concentration of 10
316 mM ($p < 0.001$, 95% CI [90.2, 205.6]) (**Figure 4**). The 95% CI represent the extent to
317 which the pH 7 and bile 10 mM conditions produce an increase in the total FFA. This is
318 in accordance to previous studies conducted by our group (Asensio-Grau et al. 2018,
319 Peinado et al. 2018)

320 Focusing on the results when considering the food products and their type of
321 matrix, more concrete patterns could be depicted (**Table 4**). Digestions conducted at
322 intestinal pH 7, revealed that in the protein fibre matrices the higher concentration of
323 bile salts led to a significant increase in total FFA release of up to 300 mg / g fat, while
324 this effect of bile concentration showed a lower increase in the unstructured matrices
325 (50 mg/ g fat, except for luncheon ham whose increase was higher). In contrast, the
326 unstructured matrices were more influenced by the 10mM bile salts concentration in the
327 intestinal pH 6 scenario, which produced an increase of around 150 mg of FFA released
328 per gram of fat as compared to the use of 1mM bile. At this last pH 6 and 1mM
329 combination, the natural protein fibres matrices did not show a common pattern.
330 Although the 10 mM concentration allowed for a higher amount of FFA release in all

331 the meat products, the increase was up to different extents: the highest was in beef steak
332 (400 mg/g fat) and the lowest in chicken drum (which showed no increase).

333 Thus, at intestinal pH 7 bile concentration plays a crucial role in natural matrices
334 while it is not relevant in the unstructured matrices; and at pH 6, the higher bile
335 concentration improves FFA release in unstructured matrices but shows a rather random
336 effect in the natural ones. On the one hand, bile salts are bio-surfactant compounds that
337 play a crucial role in lipid digestion. They adsorb onto lipid droplets and remove
338 proteins, emulsifiers and free fatty acids (lipolysis products). This mechanism facilitates
339 lipases' access and action (Maldonado-Valderrama et al. 2011, Harries et al. 1979). On
340 the other hand, the intestinal pH environment determines enzymatic activity and also
341 influences the isoelectric point of the protein that may be present at the lipid droplet
342 interface causing isoelectric effects. This fact may change the protein to a cationic form
343 allowing the bind to the anionic bile salts. Consequently, different meat matrices result
344 in different systems during digestion in which proteins can interact with the bile salts
345 either promoting or inhibiting the activity of pancreatic lipase (Bauer et al. 2005, Lowe
346 2002).

347 To sum up, unstructured matrices led to higher MDI than the structured ones;
348 higher MDI led to higher lipolysis extents (%); and the intestinal pH 7 and the 10 mM
349 bile salts concentration allowed for higher lipolysis extents. These associations,
350 explained by means of linear regression models, are summarised in **Table 5**.

351 Overall, a main limitation has been identified when aiming at discussing the
352 results. There are no studies conducted in meat products assessing lipolysis, and in
353 contrast, all of them focus on proteolysis. Nonetheless, two papers related to this topic
354 were found: one study assessing lipid digestibility of pork patties depending on the
355 thermal treatment (Hur et al. 2014), and another compiling knowledge about factors
356 affecting lipolysis in fish (Olsen and Ringø 1997). This scarcity has prevented the
357 comparison of the results to previous literature. This is a point of special relevance:
358 characterising the role of lipid digestion in meat products is of utmost necessity, as
359 recently the consumption of red meat products and cold meat derivatives has been
360 classified by the World Health Organisation as a major determinant in the development
361 of cancer (McGuire 2016). In addition, processed meat products have been traditionally
362 characterised by a high content of fat, mainly composed of saturated fatty acids. This
363 type of fat is unequivocally associated to the development of nutrition related conditions
364 such as high blood pressure and diseases like obesity (Chen et al. 2016).

365

366 **Conclusion**

367 In conclusion, fat digestibility in a wide range of meat products has been screened and
368 characterised for the first time regarding their composition, structure and intestinal
369 digestion conditions. The results reveal that lipolysis extent is dependent on the food
370 matrix degradation, and that unstructured processed meats allow for higher amount of
371 free fatty acids release than the natural protein fibrous matrices. In addition to the food
372 structure, the bile salts concentration in the digestion and the pH in the intestinal
373 digestion medium, contribute significantly to the fate of lipid digestion in meat
374 products.

375

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385

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388

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532 **TABLES**

533

534

Table 1. Composition of simulated digestion fluids.

Constituent	SSF	SGF	SIF
	mmol/ L	mmol/ L	mmol/ L
KCl	15.1	6.9	6.8
KH ₂ PO ₄	3.7	0.9	0.8
NaHCO ₃	13.6	25	85
NaCl	-	47.2	38.4
MgCl ₂ (H ₂ O) ₆	0.15	0.1	0.33
(NH ₄) ₂ CO ₃	0.06	0.5	-
CaCl ₂	1.5	0.15	0.6

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The addition of pepsin, Ca²⁺ solution and water will result in the correct electrolyte concentration in the final digestion mixture.

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SSF: Simulated Salival Fluid; SGF: Simulated Gastric Fluid; SIF: Simulated Intestinal Fluid.

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558 **Table 2.** Characterization of water, fat and protein content in the different meat matrices
 559 expressed as g/ g of product.

	Water content (g/ g product)	Fat content (g/ g product)	Protein content (g/ g product)
Hamburger	0.51 ± 0.03	0.24 ± 0.04	0.21 ± 0.02
Sausage	0.683 ± 0.002	0.107 ± 0.003	0.18 ± 0.02
Luncheon ham	0.653 ± 0.002	0.120 ± 0.006	0.11 ± 0.003
Pate	0.606 ± 0.004	0.30 ± 0.09	0.090 ± 0.002
Cured ham	0.529 ± 0.03	0.1 ± 0.02	0.32 ± 0.05
Cooked ham	0.756 ± 0.004	0.025 ± 0.03	0.199 ± 0.006
Chicken drum	0.649 ± 0.012	0.12 ± 0.03	0.201 ± 0.004
Pork loin	0.543 ± 0.026	0.135 ± 0.06	0.29 ± 0.04
Beef steak	0.56 ± 0.014	0.10 ± 0.03	0.30 ± 0.03

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580 **Table 3.** Matrix Degradation Index (%) obtained for the different meat matrices
 581 (hamburger, sausage, luncheon ham, pate, cured ham, cooked ham, chicken drum, pork
 582 loin and beef steak) after the in vitro digestion process using a fixed enzyme dose (2000
 583 LU/ g fat) and different duodenal conditions of pH and Bile concentration.

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	pH 6 - 1 mM	pH 6 - 10 mM	pH 7 - 1 mM	pH 7 - 10 mM
MDI (%)				
Hamburger	69 ± 2	69.06 ± 1.12	83 ± 4	82.1 ± 0.9
Sausage	82 ± 6	79 ± 3	94 ± 2	96.6 ± 1.2
Luncheon ham	86 ± 3	87.8 ± 0.5	90.38 ± 0.13	90.2 ± 1.4
Pate	78.6 ± 0.6	85.13 ± 1.14	81.2 ± 0.2	88.957 ± 1.014
Cured ham	70 ± 5	73.6 ± 0.2	67 ± 5	78 ± 6
Cooked ham	55.7 ± 0.3	55.13 ± 0.13	59.1 ± 1.2	63.3 ± 0.6
Chicken drum	63.57 ± 1.06	66 ± 3	69 ± 4	74 ± 2
Pork loin	52 ± 9	52 ± 6	68 ± 4	68.73 ± 0.04
Beef steak	42 ± 2	45 ± 3	51 ± 3	56 ± 2

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586 **Table 4.** Total FFA (mg/ g fat) obtained for the different meat matrices (hamburger,
 587 sausage, luncheon ham, pate, cured ham, cooked ham, chicken drum, pork loin and beef
 588 steak) after the in vitro digestion process using a fixed enzyme dose (2000 LU/ g fat)
 589 and different duodenal conditions of pH and Bile concentration.

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	pH 6 - 1 mM	pH 6 – 10 mM	pH 7 – 1 mM	pH 7 – 10 mM
Total FFA (mg/ g fat)				
Hamburger	253 ± 13	376 ± 58	501 ± 149	554 ± 115
Sausage	525 ± 22	667 ± 26	415 ± 44	490 ± 67
Luncheon ham	498 ± 18	619 ± 3	705 ± 1	902 ± 77
Pate	613 ± 153	710 ± 219	680 ± 41	700 ± 147
Cured ham	344 ± 80	467 ± 66	409 ± 85	600 ± 3
Cooked ham	298 ± 29	365 ± 162	416 ± 131	600 ± 177
Chicken drum	248 ± 8	249 ± 6	142 ± 109	408 ± 142
Pork loin	226 ± 8	614 ± 67	556 ± 104	653 ± 29
Beef steak	298 ± 4	783 ± 10	675 ± 199	860 ± 272

592 **Table 5.** Linear mixed regression models explaining the association between the study
 593 variables

Explained effect	Estimate	Standard error	95% Confidence interval (CI)	p-value
Effect of the matrix structure (unstructured vs. structured) on MDI (%)	19.38	5.818	[8.148, 30.612]	0.013
Effect of the MDI (%) on lipolysis extent (%)	8.144	0.856	[6.128, 9.606]	<0.001
Effect of the intestinal conditions on lipolysis extent (%)				
Intestinal pH (7 vs. 6)	111.19	29.47	[53.5, 168.9]	<0.001
Bile salts concentration (10 vs. 1 mM)	147.9	29.48	[90.2, 205.6]	<0.001

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595 **FIGURE LEGENDS**

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597 Figure 1. Matrix degradation index (MDI, %) of the meat products after in vitro
598 digestion with five pancreatic enzyme supplements concentrations (enzyme dose, LU/g
599 fat), classified according to the food matrix structure.

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601 Figure 2. Lipolysis extent (total FFA, mg/g fat) of the meat products after in vitro
602 digestion with five pancreatic enzyme supplements concentrations (enzyme dose, LU/g
603 fat), classified according to the food matrix structure.

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605 Figure 3. Linear mixed regression model plot. (A) Effect of the food matrix (natural or
606 processed) on matrix degradation index (MDI): processed matrices have significantly
607 higher MDI ($p = 0.013$, 95% CI [8.148, 30.612]). (B) Effect of the MDI on the total
608 FFA released: the higher the MDI, the higher the total FFA released ($p < 0.001$, 95% CI
609 [6.128, 9.606]).

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611 Figure 4. Linear mixed regression model plot. (A) Effect of the intestinal pH (6 or 7) on
612 total FFA: pH 7 allows for significantly higher amount of FFA released ($p < 0.001$, 95%
613 CI [53.466, 168.923]). (B) Effect of the bile salts concentration on the total FFA
614 released: 10mM concentration allows for significantly higher amount of FFA released
615 ($p < 0.001$, 95% CI [90.188, 205.645]).