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Title: Comparative phytochemical profile of the Elephant garlic (*Allium ampeloprasum* var. *holmense*) and the common garlic (*Allium sativum*) from the Val di Chiana area (Tuscany, Italy) before and following in vitro gastrointestinal digestion

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Abstract: This study is aimed to comparatively investigate the phytochemical profile of the common garlic (*Allium sativum* L.; CG) vs the *Allium ampeloprasum* var. *holmense*, named Elephant garlic (EG), collected in the Val di Chiana area (Tuscany, Italy), focusing on the nutritional and phytochemical properties. The results reported a lower amount of fibres in EG, underling the higher digestibility of this bulb, confirmed also to the lower sulphur containing compounds found in EG rather than in CG. Untargeted metabolomic profiling followed by supervised and unsupervised statistics allowed to depict the differences in phytochemical composition among the two bulbs, both as raw bulbs, processed following the in vitro gastrointestinal digestion process. Typical sulphur-containing compounds, such as alliin and N-gamma-glutamyl-S-allylcysteine, could notably be detected in lower amounts in EG. During the in vitro gastrointestinal digestion process, EG maintained a distinct phytochemical signature. Our findings support the distinct sensorial attributes of the bulbs.



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Dear **Editor**,

I'm sending you the manuscript "**Comparative phytochemical profile of the Elephant garlic (*Allium ampeloprasum* var. *holmense*) and the common garlic (*Allium sativum*) from the Val di Chiana area (Tuscany, Italy) before and following in vitro gastrointestinal digestion** by Ceccanti et al. for submission on the international journal Food Chemistry.

Thank you in advances for your cooperation.

Sincerely yours,

Lucia Guidi

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Highlights

- Elephant garlic (EG) had two-fold higher total phenolic content than common garlic (CG).
- Lipid-derived molecules, polyphenols and amino acids were the discriminative compounds in the raw bulbs.
- Sulphur-containing compounds were found in lower amounts in EG, underling the higher digestibility of this bulb.
- 4-hydroxybenzoic acid 4-*O*-glucoside, 5-nonadecylresorcinol and tryptophan were proposed as main biomarkers of EG consumption.

1 **Comparative phytochemical profile of the Elephant garlic (*Allium ampeloprasum* var.**
2 ***holmense*) and the common garlic (*Allium sativum*) from the Val di Chiana area (Tuscany,**
3 **Italy) before and following *in vitro* gastrointestinal digestion**

4
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35 compounds, such as alliin and N-gamma-glutamyl-S-allylcysteine, could notably be detected in
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37 distinct phytochemical signature. Our findings support the distinct sensorial attributes of the bulbs.

38

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40 polyphenols; sulphur compounds.

41 **1.Introduction**

42 The genus *Allium* includes about 700 bulbous species characterised by high diversity,
43 considering physiology and morphology aspects. Regarding that, the elephant garlic (*Allium*
44 *ampeloprasum* var. *holmense* (Mill.) Asch. et Graebn.) is classified as a type of leek (*Allium*
45 *ampeloprasum* var. *porrum* (L.) J. Gay) but, at the same time, it is considered as the common garlic
46 (*Allium sativum* L.) in terms of shape and flavour, although being three times as large as the
47 common garlic (Kim et al., 2018a). Moreover, *A. sativum* is the economically most important
48 species belonging to the *Allium* genus and has been used for long time as food and in
49 pharmacology. This bulbous species owes its importance as antifungal, antibacterial, antiviral,
50 antitoxic and anticancer agent (Rattanachaikunsopon & Phumkhachorn, 2009; Kim et al., 2018a).
51 The bioactive compounds of garlic are reported to have a biological effect on human metabolism
52 (i.e. antithrombotic and fibrinolytic effects in blood, reducing effect of LDL cholesterol level)
53 (Steiner, Khan, Holbert & Lin, 1996). These bioactive compounds of common garlic can be divided
54 into sulphur-containing compounds and sulphur-free polyphenolic compounds. The sulphur-
55 containing compounds (i.e. alliin and its derivatives) are the main responsible for the antimicrobial
56 activity, as well as of the peculiar sensorial attributes of this bulb. In contrast, the sulphur-free
57 polyphenolic compounds play an important role to prevent the oxidative damage caused by ROS
58 (Reactive Oxygen Species) (Ma et al., 2011).

59 Nowadays, elephant garlic has been proposed as a substitute of common garlic in cooking (fresh
60 or processed) because its flavour is very close to that of common garlic but with a milder impact on
61 human breath and a better digestibility than common garlic (Block, 2011). For this reason, the
62 elephant garlic is named “kissingarlic”, “garlic for people who don’t like garlic” and “garlic-like”
63 (Lu, Ross, Powers, Aston & Rasco, 2011). In Val di Chiana, an area located in Tuscany (centre of
64 Italy), with peculiar weather and soil characteristics, elephant garlic was joined to the list of
65 Traditional Agri-food Products of the Tuscany Region (Executive Decree Tuscany Region, n. 1569
66 of April 4th, 2016

67 https://www.aglionevaldichiana.net/public/Documenti/Decreto_Regione_Toscana.pdf) and later to
68 the list of Traditional Agri-food Products of Italy (G.U. n.143 of June 21th, 2016
69 https://www.aglionevaldichiana.net/public/Documenti/Decreto_MiPAAF.pdf) with the name
70 “Aglione della Valdichiana”.

71 In nutritional and nutraceutical terms, the elephant garlic features suggest a reduction in the
72 content of sulphur-compounds among its bioactive substances, although few studies have been
73 carried out for its characterization to date. Kim et al. (2018b) evaluated the organo-sulphur
74 compounds in *Allium* species, showing a high content of γ -glutamyl peptides in the elephant garlic
75 and the highest alliin content in the common garlic. Therefore, they reported a higher content of
76 sulphur-containing volatile compounds in common garlic rather than in the elephant garlic, with the
77 presence of 13 sulphur compounds found in garlic against 6 sulphur compounds found in the
78 elephant garlic, even though in very low concentrations. Moreover, Najda, Błaszczyk, Winiarczyk,
79 Dyduch and Tchórzewska (2016), analysing elephant garlic and common garlic from Poland, found
80 a higher polyphenol content and antioxidant activity in elephant garlic bulb rather than in common
81 garlic bulb. These authors did not find peculiar differences in terms of polyphenol profile between
82 the two bulbs. In contrast, Lu, Ross, Powers, Aston and Rasco (2011), analysing American garlics
83 and elephant garlics, found lower antioxidant activity in the elephant garlic rather than in the
84 common garlic and a polyphenol content very similar in both the bulbs. The comparison of these
85 last two studies suggests that the origin of both the bulbs may play a pivotal role in determining the
86 polyphenol profile.

87 Holistic approaches like metabolomics could be very useful to describe and to differentiate the
88 profile of bioactive compounds in garlic and elephant garlic, as well as to ensure geographical
89 traceability (Maietti et al., 2012). This last aspect becomes relevant with the view of a hypothetic
90 award of PDO (Protected Designation of Origin), as proposed for “Aglione della Valdichiana”. In
91 fact, the quali- quantitative differences in secondary metabolites, such as sulphur compounds and
92 polyphenols, can better discriminate the bulbs hence identifying potential counterfeits thus ensuring

93 traceability. Noteworthy, a deep profiling of the phytochemicals in these two bulbs opens the
94 possibility to further investigate nutraceutical properties rather than desired or unpleasant sensorial
95 attributes.

96 In this regard, combining *in vitro* gastrointestinal digestion process with untargeted
97 metabolomics may be useful to analyse the bioaccessibility of polyphenols and other health-related
98 compounds. To date, limited information on the changes of bioactive compounds in garlic and/or
99 elephant garlic during simulated gastrointestinal processes is present in literature (Bhatt & Patel,
100 2013; Torres-Palazzolo, Ramirez, Locatelli, Manucha, Castro & Camargo, 2018; Rosen et al.,
101 2001). In the last few years, many research reported the bioaccessibility of health-promoting
102 compounds, considering it as the percentage of compounds from the food sample released during
103 the simulation of digestion (Pérez-Vicente, Gil-Izquierdo, García-Viguera, 2002; Rodríguez-Roque,
104 Rojas-Graü, Elez-Martínez & Martín-Belloso, 2014; Rocchetti, Chiodelli, Giuberti & Lucini, 2018).
105 The combination of *in vitro* gastrointestinal digestion with untargeted metabolomics may provide a
106 better understanding of the main changes occurring to bioactive compounds during simulated
107 gastrointestinal processes (Rocchetti, Chiodelli, Giuberti & Lucini, 2018).

108 Up to now, no detailed comparative studies of nutritional parameters and nutraceutical
109 compounds and their bioaccessibility during the human digestion of the elephant garlic and the
110 common garlic have been carried out. In this study, for the first time, the nutritional aspects, some
111 mineral elements (including the sulphur) as well as phenols and secondary metabolites from
112 untargeted metabolomics have been comparatively investigated in elephant and common garlic
113 offered by Val di Chiana farmers. Then, the fate of garlic metabolites was investigated for the first
114 time, using an *in vitro* gastrointestinal digestion and then an untargeted metabolomics-based
115 approach, in order to further explore the main differences between the two bulbs from a nutritional
116 standpoint.

117 **2. Materials and methods**

118 *2.1. Plant material*

119 Common garlic (CG) and elephant garlic (EG) samples were grown in Val di Chiana area and
120 were offered by local association “Qualità e Sviluppo Rurale srl” to the Department of Agriculture,
121 Food and Environment (DAFE) of the University of Pisa during June 2019. Both the bulbs derived
122 from the same farm and have been cultivated in the same climatic and edaphic conditions to
123 exclude the contribution of pedo-climatic conditions. Cloves of EG and CG were randomly
124 selected, manually peeled and chopped in small pieces. A part of the bulb material was lyophilized
125 for the determination of some mineral elements, whereas the fibre, the sugar contents and the
126 untargeted metabolomics based on high-resolution mass spectrometry; the other part was freeze-
127 dried and stored at $-80\text{ }^{\circ}\text{C}$ until analysis. All the analyses were carried out in triplicate.

128 *2.2. Proximate analysis*

129 A part of freeze-dried material was weighed and oven-dried at $65\text{ }^{\circ}\text{C}$ till constant weight and the
130 percentage of dry matter (% DM) was calculated. Then, the dried samples were used for the
131 determination of phosphorous, potassium, calcium and magnesium concentration in both the
132 analysed bulbs. Dry tissues were mineralized for 60 min at $220\text{ }^{\circ}\text{C}$ using a solution of $\text{HNO}_3:\text{HClO}_4$
133 (2.5:1 v/v). Phosphorus concentration was determined colorimetrically using an Ultrospec 2100 Pro
134 spectrophotometer (GE Healthcare Ltd., Little Chalfont, UK), following the Olsen method, whereas
135 K, Ca and Mg with an atomic absorption spectrophotometer (Varian AA 24FS, Australia). Results
136 were expressed as % P, K, Ca and Mg. Nitrogen, carbon, hydrogen and sulphur determinations
137 were obtained from an Elementar Vario MICRO cube instrument.

138 Sugar (sucrose and glucose) quantification was carried out according to Yusof, Rasmusson and
139 Galindo (2016) and Sotelo, Pérez, Najjar-Rodriguez, Walter and Dorn (2014) with minor
140 modifications. Sucrose and glucose were determined using K-SUFRG commercial kit (Megazyme,
141 Wicklow, Ireland), following the manufacturer’s protocol. Results were expressed as $\text{g}\cdot 100\text{ g}^{-1}$ dry
142 weight (DW).

143 Protein determination was performed using the spectrophotometer and the Protein Assay Kit II®
144 (Bio- Rad). Using a bovine serum albumin standard curve, the results were expressed as mg protein
145 per g fresh weight (FW).

146 The crude fibre and the fibre fractions [neutral detergent fibre (NDF) and acid detergent fibre
147 (ADF), acid detergent lignin (ADL), hemicellulose and cellulose] were analysed according to the
148 method described by Van Soest, Robertson and Lewis (1991) using the instrument ANKOM
149 (ANKOM 65 rpm agitation). Results were expressed as percentage (%).

150 *2.3. Total phenolic content*

151 For total phenolic extraction, the fresh material (1 g) was finely ground in a mortar, suspended in
152 4 mL 80% aqueous methanol (v/v), and placed in an ultrasonic water bath (Digital ultrasonic
153 Cleaner, DU-45, Argo-Lab, Modena, Italy) at 4 °C for 30 min. For the determination of total
154 phenolic content, the solution was centrifuged at 10,000 g for 7 min and an amount of the
155 supernatant was added to Folin-Ciocalteu reagent, following Folin-Ciocalteu method described by
156 Dewanto, Wu, Adom and Liu (2002) with slight modifications. Consequently, 1.25 mL Na₂CO₃ 7%
157 (w/v) were added to the solution and samples were incubated for 90 min in dark conditions. The
158 increase of absorbance was measured spectrophotometrically at 760 nm wavelength against a blank.
159 The total phenolic content was expressed as milligrams equivalents of gallic acid per g of fresh
160 weight (mg GAE g⁻¹ FW).

161 *2.4. Extraction and untargeted metabolomic profiling by UHPLC-QTOF mass spectrometry*

162 One gram of each lyophilized samples was homogenized in 10 mL of 0.1% HCOOH in 80%
163 (v/v) methanol solution using a homogenizer-assisted extraction system (Ultra-Turrax, IkaT25,
164 Staufen, Germany) as previously reported (Rocchetti, Bhumireddy, Giuberti, Mandal, Lucini &
165 Wishart, 2019). Afterwards, samples were centrifuged at 6,000 g for 15 min at 4 °C and then
166 supernatants were filtered with 0.22 µm cellulose syringe filters in vials which were stored at -18
167 °C until analysis.

168 Subsequently, the untargeted metabolomic profile of both bulbs was investigated using UHPLC-
169 ESI-QTOF mass spectrometry, as previously described (Rocchetti, Bhumireddy, Giuberti, Mandal,
170 Lucini & Wishart, 2019). Briefly, chromatography was carried out in the reverse phase mode using
171 an Agilent Zorbax eclipse plus C18 column and a water-acetonitrile gradient solution (from 6% up
172 to 90% acetonitrile in 33 min) for separation. For mass spectrometry detection, QTOF was operated
173 in positive scan mode to acquire ions in the range 50–1200 m/z . A volume of 6 μL of each extracted
174 sample was injected using nitrogen as both sheath gas (10 L min^{-1} at 350 $^{\circ}\text{C}$) and drying gas (8 L
175 min^{-1} at 330 $^{\circ}\text{C}$). The annotation of garlic and elephant garlic metabolites was achieved using the
176 software Profinder B.07 from Agilent Technologies, according to the “find-by-formula” algorithm.
177 In particular, the annotations were recursively achieved against the comprehensive database
178 FoodDB (www.fooddb.ca), one of the most comprehensive databases available in literature for
179 untargeted studies in food metabolomics and using the entire isotopic profile with a maximum of 5
180 ppm for mass accuracy. Therefore, in our experimental conditions, a Level 2 of compound
181 identification was achieved as set out by the COSMOS Metabolomics Standards Initiative
182 (Rocchetti, Giuberti, Busconi, Marocco, Trevisan, & Lucini, 2020; Salek, Neumann, Schober,
183 Hummel, Billiau & Steinbeck, 2015; Schrimpe-Rutledge, Codreanu, Sherrod & McLean, 2016).
184 The obtained dataset was further used for statistics and chemometrics.

185 2.5. Simulated *in vitro* gastrointestinal digestion process

186 The *in vitro* gastrointestinal digestion, simulating the oral, gastric and pancreatic digestion
187 phases, was applied to lyophilized samples according to the static method detailed by Minekus et al.
188 (2014). Samples (250 mg) were homogenized with 175 μL simulated salivary fluid (SSF) (Minekus
189 et al., 2014) at pH 7.0, 25 μL salivary α -amylase (from human saliva Type IX-A, Sigma) solution
190 made up in SSF electrolyte stock solution, 1.25 μL CaCl_2 0.3 M and 48.75 μL water. The oral step
191 was run at 37 $^{\circ}\text{C}$ for 2 min. Then, the oral bolus samples were mixed (ratio 1:1) with 375 μL
192 simulated gastric fluid (SGF) at pH 3.0, 80 μL porcine pepsin stock solution (25,000 U mL^{-1}) made
193 up in SGF electrolyte stock solution (pepsin from porcine gastric mucosa, Sigma), 0.25 μL CaCl_2 0.3

194 M, 10 μ L HCl 1 M to adjust pH to 3.0, and rest of volume with water. The gastric phase was
195 carried out for 2 h at 37 °C. Then, gastric chyme was mixed (1:1) with 550 μ L simulated intestinal
196 fluid (SIF) electrolyte stock solution consisting at pH 7.0, 250 μ L of a pancreatin solution (800 U
197 ml^{-1}) based on pancreatin α -amylase activity made up in SIF electrolyte stock solution (pancreatin
198 from porcine pancreas, Sigma), 125 μ L fresh bile (160 mM in fresh bile), 2 μ L CaCl_2 0.3 M, 7.5 μ L
199 NaOH 1 M to adjust pH to 7.0, and water filling the rest of the volume. The intestinal phase was
200 carried out for 2 h at 37 °C. At selected time points (i.e., gastric and pancreatic phases)
201 corresponding digestion sample tubes for each material were cooled on ice to stop the reaction. The
202 experiment was performed in triplicate. Finally, to depict the fate of bioactive compounds during
203 the *in vitro* gastrointestinal digestion, digested samples were prepared and analysed using an
204 untargeted UHPLC-ESI/QTOF mass spectrometry as above reported for the undigested materials.

205 2.6. Statistical analysis and chemometrics

206 Results of proximate analysis, minerals and total phenolic content were compared with Student *t*-
207 test ($P \leq 0.05$). Data are expressed as mean \pm standard deviation. This statistical analysis was
208 performed using GraphPad (GraphPad, La Jolla, CA, USA).

209 Metabolomic data were interpreted using Agilent Mass Profiler Professional B.12.06 (from
210 Agilent Technologies). Compounds were filtered by abundance and by frequency (only those
211 compounds with an area > 5000 counts and appearing in 100% of samples in at least one condition
212 were considered), normalized at the 75th percentile and baselined to the median of each compound
213 in all samples. The unsupervised hierarchical cluster analysis (HCA – Euclidean distance) was then
214 used to naively group samples, according to intrinsic similarities in metabolomic profile (Rocchetti,
215 Bhumireddy, Giuberti, Mandal, Lucini & Wishart, 2019). Afterwards, the dataset was exported into
216 SIMCA 13 (Umetrics, Malmo, Sweden), Pareto scaled and elaborated for orthogonal partial least
217 squared discriminant analysis (OPLS-DA) supervised modelling, considering the combination of
218 "sample type x digestion phase" as class membership criterion. Finally, the variable importance in
219 projection (VIP analysis) method was used to evaluate the discrimination potential of the different

220 metabolites (i.e., those compounds possessing a VIP score > 1) and a Fold-Change analysis (FC >
221 5) was combined with ANOVA (P < 0.01, Bonferroni multiple testing correction) in Volcano plot
222 to point out differential metabolites between raw common garlic and elephant garlic samples.

223 **3.Results and discussion**

224 *3.1. Nutritional results*

225 Results of the proximate analysis of CG and EG cloves cultivated in Val di Chiana area were
226 reported in Table 1. No differences were found in sugars (glucose and fructose) content of both the
227 analysed bulbs, whereas a higher protein content was recorded in EG than in CG. Even the dry
228 matter and the moisture contents were different comparing both the bulbs, with EG reporting a
229 higher moisture content than CG (Table 1). NDF and cellulose resulted significantly lower in the
230 EG as compared to CG, underling the superior digestibility of this bulb (Baer, Rumpler, Miles &
231 Fahei, 1997). These differences probably reflect the differences in variety and species among the
232 two analysed bulbs as well as the different conditions of growth. Mineral content in the bulb
233 samples differed significantly for P and K exclusively and these minerals were lower in the EG as
234 compared to the CG (Table 1). An interesting result is the higher content of the sulphur element
235 found in EG (+ 41%) as compared with CG. Mineral results are comparable with the findings in
236 garlic of Sajid, Butt, Shehzad and Tanweer (2014) and Odebunmi, Oluwaniyi and Bashiru (2010).

237 *3.2. Phytochemical discrimination of raw samples*

238 The first result that need to be highlighted in terms of functional components is represented by
239 the total phenolic content, which was found significantly higher in EG than in CG (Fig. 1). This
240 result is in agreement with the findings by Lu, Ross, Powers, Aston and Rasco, (2011) whilst
241 contrasting outcomes that have been provided by Najda, Błaszczuk, Winiarczyk, Dyduch and
242 Tchórzewska (2016), which reported very low total phenolic content in both bulb types. Thereafter,
243 untargeted metabolomics based on UHPLC-QTOF mass spectrometry was used to investigate in a
244 comprehensive way the differences and similarities in the phytochemical of the two bulb samples,
245 both before and after the *in vitro* gastrointestinal digestion process. Overall, this approach allowed

246 to putatively annotate 2745 mass features that were classified according to the database FoodDB,
247 together with individual abundances and composite mass spectra (Table S1). As first evaluation, a
248 Volcano plot was produced to compare raw EG vs CG by coupling ANOVA ($P \leq 0.01$) and Fold-
249 Change (cut-off ≥ 5) analysis (Table S2). As it can be observed, 161 metabolite species were found
250 to discriminate EG and CG, thus suggesting distinctive chemical fingerprints of the raw matrices
251 before *in vitro* gastrointestinal digestion. Interestingly, only the 22% of the discriminant markers
252 were found to be down-accumulated in EG when compared with CG; among these compounds, we
253 found typical compounds characterizing garlic such as alliin (belonging to α -amino acids), together
254 with two isomeric dipeptides, namely N-gamma-glutamyl-S-allylcysteine and N-gamma-glutamyl-
255 S-cis-(1-propenyl)cysteine. The down-accumulation of sulphur, containing compounds such as N-
256 gamma-glutamyl-S-allylcysteine and N-gamma-glutamyl-S-cis-(1-propenyl)cysteine, characterized
257 EG (Table S2) and could explain the higher total phenolic content observed in EG compared to CG
258 (Fig. 1) since Phan, Netzel, Chhim, Netzel and Sultanbawa (2019) reported that the total phenolic
259 content can decrease with the increase in organosulfur compounds and terpenoid substances in
260 mature garlic bulbs. In addition, a significant down-accumulation of 6 polyphenols and 8 prenol
261 lipids was outlined in EG when compared with CG (Table S2).

262 Overall, it has been suggested that the biological and health promoting properties of garlic
263 primarily derived from its polyphenols and organosulfur compounds (Phan, Netzel, Chhim, Netzel
264 & Sultanbawa, 2019). These trends were confirmed by Kim et al. (2018b), reporting a high level of
265 bioactive γ -glutamyl peptides in both EG and CG. However, considering that organosulfur
266 compounds are extremely unstable and susceptible to further transformation into volatile
267 compounds (such as allicin and diallyl-sulfides), recent attention has been placed on polyphenols
268 due to their potential role in health-related benefits to humans. On the other hand, 125 additional
269 compounds (i.e., the remaining 78% of discriminant metabolites outlined by Volcano plot analysis)
270 were proposed in this work as chemical markers of EG. The most represented classes among the
271 discriminant markers were those of steroids and derivatives (27 compounds), glycerol- and

272 glycerophospho- lipids (24 compounds), prenol lipids (13 compounds), polyphenols (12
273 compounds), amino acids and derivatives (8 compounds), organooxygen compounds (8
274 annotations) and fatty acyls (7 compounds). Therefore, our findings revealed a large presence of
275 compounds belonging to lipids and steroids (mainly saponins) in EG. These results are not
276 surprising; in fact, plants belonging to the genus *Allium* have been previously reported as a good
277 source of bioactive saponin compounds, responsible for many of their reported pharmacological
278 activities (e.g., antiproliferative, antifungal and antispasmodic activities). In this regard, previous
279 studies (Lanzotti, 2005; Petropoulos, Fernandes, Ntatsi, Petrotos, Barros & Ferreira, 2018) showed
280 that saponins characterized by furostane, spirostane, cholestane, and oleanane-type structures are
281 widely represented in *Allium*, thus confirming our findings.

282 Additionally, an abundance of N-gamma-L-glutamyl-L-methionine and eruboside B was noticed
283 in the EG sample (Table S1). N-gamma-L-glutamyl-L-methionine belongs to the organosulphur
284 compounds with important biological effects (lipid-lowering, antidiabetic, anticancer, anti-
285 asthmatic, antiplatelet and anti-atherosclerotic activities) already described by Kim et al. (2018b),
286 while eruboside B is a typical garlic compound that improve the antimicrobial properties of *Allium*
287 vegetables (Nakamoto, Kunimura, Suzuki & Kodera, 2020). However, it is important to take into
288 account that genotype has a great impact on the metabolomic profile of garlic and elephant garlic
289 bulbs (Najda, Błaszczuk, Winiarczyk, Dyduch & Tchórzewska, 2016); therefore, both genotype and
290 pedoclimatic conditions represent two critical parameters that need to be taken always into account
291 to reach the quality improvement of the final products.

292 3.3. *In vitro* gastrointestinal digestion and discrimination of both bulb samples

293 Moreover, once the differences between EG and CG were represented in the raw matrices,
294 multivariate statistics (based on both unsupervised and supervised methods) were used to depict the
295 changes occurring during the *in vitro* gastrointestinal digestion process of both bulbs. On this
296 matter, unsupervised hierarchical cluster analysis (i.e., HCA) carried out on the UHPLC-QTOF
297 mass spectrometry data allowed to identify a clear separation trends (Fig. 2), outlining a strong

298 impact of each digestion phase (i.e., both gastric and pancreatic phases) on the phytochemical
299 composition of both bulb samples. Initial differences in phytochemical profiles between EG and CG
300 may be notably conserved, even during the digestion process. The results from unsupervised
301 statistics suggest that a further application of the OPLS-DA score plot would help to point out the
302 most discriminant compounds, driving the trends observed. The OPLS-DA score plot illustrating
303 modification of the metabolite profiles, moves from raw samples to digested samples and it is
304 provided in Fig. 3. This supervised model allowed to confirm the unsupervised findings (Fig. 2).
305 Firstly, a confirmation of the differences existing on the raw matrices was noticed on the left part of
306 the graph, confirming the results of Volcano plots previously discussed (Table S2). Besides, the
307 second latent vector $t[2]$ showed a clear impact of both gastric and pancreatic phases of digestion,
308 driven also by the different matrix incubated (i.e., EG vs CG) (Fig. 3).

309 Furthermore, a variable selection method (VIP; variable importance in projection) was used to
310 reduce the numbers of variables and better explain the differences observed over the *in vitro*
311 digestion process (Table 2). Markers, that assigned a VIP score > 1 , are organized in chemical class
312 and reported together with the Log Fold-Change (FC) values for each main comparison (i.e., EG vs
313 CG on raw, gastric and pancreatic samples) (Table 2). Overall, 85 compounds matched this
314 criterion (including some isomeric compounds), mainly belonging to the classes of polyphenols,
315 amino acids, benzenoids, sulphur containing compounds, fatty acyls, glycerophospholipids,
316 heteroaromatic compounds, indoles, prenol lipids, pyrrolizines, quinolines, steroids and derivatives,
317 tetrahydrofurans and other compounds. Overall, when considering the comparison EG vs CG during
318 the gastric phase of digestion, the most affected compounds were found to be hovenidulcigenin A (a
319 prenol lipid; LogFC = -18.55), LysoPC(18:1(11Z)) (a glycerophospholipid; LogFC = -18.36) and
320 4-[(2-hydroxy-1-naphthalenyl)azo]benzenesulfonic acid (a benzenoid; LogFC = -9.28). Until
321 today, the bioactive and pharmacological role of the hovenidulcigenin A and 4-[(2-hydroxy-1-
322 naphthalenyl)azo]benzenesulfonic acid has not been established in literature, while the LysoPC
323 species play an important role as lipid mediators in cellular responses and pathophysiology; in fact,

324 they are involved in the activation of inflammatory responses and their potential role as vaccine has
325 been discussed (Wi, Seo, Cho, Nam & Park, 2014).

326 Moreover, the gastric phase of digestion mainly affected CG polyphenols composition. In fact,
327 the discriminant compounds, namely 4-hydroxybenzoic acid 4-*O*-glucoside, phloretin and
328 butein/naringenin were characterized in EG during the gastric phase by LogFC values > 2. The 4-
329 hydroxybenzoic acid is a phenolic acid which can be converted into more useful compounds such as
330 resveratrol, muconic acid, gastrodin, ubiquinone with a wide variety of biological and
331 pharmaceutical activities as antibacterial, antioxidant, anticancer, hypolipidemic, prevention of
332 heart diseases activities (Wang, Bilal, Hu, Wang, & Zhang, 2018); phloretin is one of the best
333 known dihydrochalcone with antifungal, antiviral, anti-inflammatory, estrogenic, anticancer and
334 estrogenic activity and able to improve the fluidity of biological membranes, increasing the
335 penetration of drugs (Behzad, Sureda, Barreca, Nabavi, Rastrelli, & Nabavi, 2017) and
336 butein/naringenin are flavonoids with important healthy roles such as antioxidant, antitumor,
337 cardioprotective, antiviral and antibacterial activity (Salehei et al., 2019; Bordoloi et al., 2019).

338 Finally, analysing the results of the pancreatic phase, EG was characterized by an overall down-
339 accumulation of several lipid-derived compounds, mainly belonging to fatty acyls (6 annotated
340 compounds) and steroids (4 annotated compounds). Afterwards, the polyphenol 4-hydroxybenzoic
341 acid 4-*O*-glucoside showed significantly higher LogFC values (i.e., 9.95) in EG when compared to
342 CG also during pancreatic digestion phase. Other metabolites characterizing EG during pancreatic
343 phase of digestion were pantoyllactone glucoside (fatty acyls), (2*S*,4*S*)-monatin (alpha amino acid),
344 followed by several isomeric compounds classified as "other compounds" (Table 2). The OPLS-DA
345 model allowed to detect also discriminant compounds characterizing the raw bulbs; for example,
346 AS 1-5 (belonging to the class of organic compounds known as glycosyl-*n*-acylsphingosines and
347 typical in garlic) was found to characterized raw EG (LogFC = 19.91) sample, but it was heavily
348 affected by the *in vitro* digestion process (Table 2).

349 Regarding the potential biomarkers proposed in this work, 4-hydroxybenzoic acid 4-*O*-
350 glucoside, 5-nonadecylresorcinol and tryptophan were found to be among the most representative
351 compounds in the undigested EG (Table 2). According to literature, phenolic acids such as 4-
352 hydroxybenzoic acid 4-*O*-glucoside can be released from the food matrix in the stomach, further
353 enhancing their release and absorption. The absorption of phenolic acids from beverages occurs at a
354 higher extent than from solid food matrices. In fact, most of phenolic acids exist as conjugated or
355 bound to dietary fibre, thus reaching the colon and becoming available for further metabolism by
356 gut microbiota (Rocchetti et al., 2019; Mosele, Maciá & Motilva, 2015). In addition, hydrolysis by
357 intestinal or microbial esterases can promote the release of phenolic acids in the intestine,
358 supporting their absorption across the gastrointestinal barrier and enter the peripheral blood
359 circulation. Therefore, besides a clear species effect, our findings suggested that consuming EG
360 could be a valid strategy to promote the bioaccessibility of bioactive phenolic acids, such as 4-
361 hydroxybenzoic acid 4-*O*-glucoside. Another phenolic compound namely 5-nonadecylresorcinol is
362 an alkylresorcinol with a cytotoxic activity, and it is able to significantly inhibit the growth of
363 various cell lines, such as lung cancer cells, human epithelial cells, breast cancer cells, epithelioid
364 cervix carcinoma cells and human central nervous system tumour cell line (Liu, Winter, Stevenson,
365 Morris & Leach, 2012). The tryptophan is an essential amino acid for the human health because of
366 human body is not able to synthesize it. Recently, Li et al., (2019) reported a high antioxidant
367 activity of this amino acid and the addition of tryptophan to the walnut protein-derived peptides
368 had a potential in the inhibition of xanthine oxidase, a critical enzyme in human health, because of
369 its ability to catalyse the oxidation of hypoxanthine to xanthine and xanthine to uric acid. Therefore,
370 the inhibition of the xanthine oxidase may be able to alleviate the development of hyperuricemia (Li
371 et al., 2019). The different accumulation of health-related compounds in both analysed bulbs,
372 during the *in vitro* gastrointestinal digestion process, supports the potential exploitation of EG as
373 source of bioactive compounds with important biological and pharmacological roles in addition to
374 CG.

375 **Conclusions**

376 This study is aimed at comparing phytochemical compounds of CG and EG from the Val di
377 Chiana area (Tuscany, Italy), with a focus on bioactive compounds. In fact, despite these two bulbs
378 could seem similar in shape and aspect, they belong to different *Allium* species. The differences
379 were reflected also in the proximate results and produced two distinctive metabolomic profiles.
380 Firstly, the EG samples showed a lower fibre content, which is supportive for a higher digestibility
381 of this bulb, confirmed successively to the lower sulphur-containing compounds found in EG rather
382 than in CG. The total phenolic content definitely resulted two-fold higher in the EG than in the CG.
383 Finally, the untargeted metabolomic approach using UHPLC-QTOF mass spectrometry allowed to
384 identify a higher number of organosulphur compounds in CG than in EG. These sulphur-containing
385 metabolites are responsible for several biological effects of *Allium* vegetables, in addition to some
386 unpleasant attributes that garlic leaves in breath of humans. This opens the possibility to use EG to
387 replace CG with food tasting purposes. The untargeted metabolomic approach also identified 125
388 key metabolites which were most representative in raw EG, mainly including lipid-derived
389 molecules, polyphenols and amino acid derived compounds. In addition, clear differences were
390 outlined between EG and CG during *in vitro* gastrointestinal digestion process, underling a higher
391 impact of the gastric phases on the phytochemical modifications, with 4-hydroxybenzoic acid 4-*O*-
392 glucoside, 5-nonadecylresorcinol and tryptophan proposed as biomarker of the consumption of EG.
393 Taken together, the present findings indicate distinct phytochemical profiles among EG and CG,
394 with distinct bioactive and functional properties and sensorial attributes. Our dataset also
395 contributes to identify some putative biomarkers that could be exploited for the traceability of
396 *Allium ampeloprasum* var. *holmense*.

397

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402 The authors declare that they have no known competing financial interests or personal relationships
403 that could have appeared to influence the work reported in this paper.

404 **CRedit authorship contribution statement**

405 **Costanza Ceccanti:** Conceptualization, Data curation, Formal analysis, Writing - original draft.

406 **Gabriele Rocchetti:** Conceptualization, Methodology, Investigation, Formal analysis, Writing -

407 review & editing. **Marco Landi:** Conceptualization - review & editing, Supervision. **Gianluca**

408 **Giuberti:** Methodology, Investigation, Formal analysis, Writing- review & editing. **Luigi Lucini:**

409 Validation - review & editing, Supervision. **Stefano Biagiotti:** Resources, Supervision. **Lucia**

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414 **Appendix A. Supplementary data**

415 The following are the Supplementary data to this article:

416 • Table S1

417 • Table S2

418 **References**

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Figure Captions

Figure 1. Total phenolic content of common garlic and elephant garlic. Data were compared with Student *t*-test ($P \leq 0.05$). Significance ***: $P \leq 0.001$.

Figure 2. Unsupervised hierarchical cluster analysis (HCA) based on Fold-Change heat map (similarity: Euclidean; linkage rule: ward) for raw and *in vitro* digested elephant garlic (EG) and common garlic (CG) samples.

Figure 3. Supervised OPLS-DA prediction model for raw and *in vitro* digested elephant garlic (EG) and common garlic (CG) samples.

562

563 **Table 1.** Proximate analysis and minerals of edible garlic and elephant garlic cloves. Data were compared with Student *t*-test ($P \leq 0.05$).564 Significance ns: not significant; *: $P \leq 0.05$; **: $P \leq 0.01$; ***: $P \leq 0.001$ for the interaction of factors.

Proximate composition		Unit	Garlic (<i>Allium sativum</i>)	Elephant garlic (<i>Allium ampeloprasum</i> var. <i>holmense</i>)	Significance
Protein		g 100 g ⁻¹ FW	0.98±0.05	1.22 ± 0,26	*
Carbohydrate	Glucose	g 100 g ⁻¹ DW	3.55±0.52	3.36 ± 0.14	ns
	Sucrose		0.11±0.04	0.11 ± 0.03	ns
Dietary fibre	Neutral detergent fibre (NDF)		9.98±0.16	8.46 ± 0.30	**
	Acid detergent fibre (ADF)		9.09±0.40	8.58 ± 1.09	ns
	Acid detergent lignin (ADL)	%	3.91±0.34	3.25 ± 0.31	ns
	Hemicellulose		0.48±0.33	0.55 ± 0.12	ns
	Cellulose		2.06 ± 0.23	1.29 ± 0.04	**
Minerals	Nitrogen (N)		2.64 ± 0.26	1.88 ± 0.06	**
	Carbon (C)		41.29 ± 0.26	42.38 ± 0.28	**
	Hydrogen (H)		6.56 ± 0.02	6.68 ± 0.04	*
	Sulphur (S)	%	0.38 ± 0.01	0.65 ± 0.18	*
	Phosphorus (P)		1.17 ± 0.04	0.59 ± 0.01	***
	Potassium (K)		1.04 ± 0.11	0.76 ± 0.02	*
	Calcium (Ca)		0.21 ± 0.01	0.23 ± 0.02	ns
	Magnesium (Mg)		0.05 ± <0.01	0.06 ± <0.01	ns
Moisture content		%	62.13 ± 0.57	67.23 ± 0.67	***
Dry matter		%	37.87 ± 0.57	32.77 ± 0.67	***

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567 **Table 2.** Bioactive compounds identified by VIP (Variable Importance in Projection) selection method following OPLS-DA model, in the raw
 568 matrix of elephant garlic and common garlic and during the *in vitro* gastrointestinal digestion of the two analysed bulbs. Compounds are provided
 569 together with VIP scores (measure of variable's importance in the OPLS-DA model) and Log Fold-Change in raw material and during different
 570 phases (gastric and pancreatic phases) of the *in vitro* digestion. ¹ns: not significative.

Class	Metabolites	VIP score (OPLS-DA)	LogFC [EG vs CG] raw	LogFC [EG vs CG] gastric	LogFC [EG vs CG] pancreatic
<i>Polyphenols</i>	4-Hydroxybenzoic acid 4-O-glucoside	1.02±0.56	11.42	2.16	9.95
	Rosmarinic acid	1.06±0.25	-18.70	-0.83	ns
	Sinapoylspermine	1.22±0.23	¹ ns	-0.37	-17.73
	Bisdemethoxycurcumin	1.22±0.27	0.35	0.64	2.31
	5-Nonadecylresorcinol	1.00±0.53	21.89	ns	ns
	Phloretin	1.06±0.14	0.49	9.80	ns
	Butein/Naringenin	1.24±0.32	ns	11.23	0.01
	<i>Aminoacids and derivatives</i>	L-Proline	1.02±0.58	-2.56	-1.01
N-acetyl lysine methyl ester		1.23±0.33	9.24	9.28	-0.33
Tryptophan		1.23±0.35	19.61	-0.46	-0.29
Cinnamoylglycine		1.22±0.36	0.57	1.08	-0.30
<i>Benzenoids</i>	(2S,4S)-Monatin	1.22±0.33	0.10	0.64	19.61
	N-Acetylarylamine/ N-benzylformamide/2-Phenylacetamide	1.22±0.32	17.95	1.23	-0.31
	2-(2-Methylpropoxy)naphthalene	1.23±0.33	ns	9.28	-0.33
	4-[(2-Hydroxy-1-naphthalenyl)azo]benzenesulfonic acid	1.23±0.11	0.56	-9.28	ns
	1,1'-[1,12-	1.24±0.35	ns	16.89	0.03

<i>Sulphur containing compounds</i>	Dodecanediylbis(oxy)]bisbenzene N-gamma-glutamyl-S-allylcysteine/ N-gamma-glutamyl-S-trans-(1-propenyl)cysteine	1.01±0.38	-7.01	0.64	2.31
<i>Fatty acyls</i>	Methyl (R)-3-methyl-2-oxopentanoate Pantoyllactone glucoside Sativic acid/ Pinellic acid/9,12,13-TriHOME/(9S,10E,12S,13S)-9,12,13-Trihydroxy-10-octadecenoic acid/5,8,12-Trihydroxy-9-octadecenoic acid/9,10,13-TriHOME Cervonoyl ethanolamide	1.22±0.19 1.23±0.27 1.23±0.13 1.24±0.38	ns 0.23 ns -7.76	0.17 0.57 -0.01 7.80	20.27 18.85 -19.08 0.16
<i>Glycerophospholipids</i>	LysoPC(18:1(11Z)) LysoPC(20:3(8Z,11Z,14Z)) LysoPC(20:4(8Z,11Z,14Z,17Z)) 1-16:0-2-18:1-phosphatidylcholine 1-18:3-2-18:1-phosphatidylcholine PE-NMe2(16:0/16:0)	1.24±0.36 1.24±0.39 1.24±0.37 1.25±0.16 1.25±0.18 1.25±0.32	-0.65 ns ns 0.41 0.34 0.15	-18.36 -1.32 -1.32 17.52 18.55 -1.01	-0.15 0.15 -0.01 ns 0.16 ns
<i>Heteroaromatic compounds</i>	5-(2-Furanyl)-3,4-dihydro-2H-pyrrole 3,4-Dihydro-4-[(5-methyl-2-furanyl)methylene]-2H-pyrrole	1.22±0.32 1.25±0.19	17.96 20.92	1.23 0.18	-0.31 0.09
<i>Indoles and derivatives</i>	3-(1H-Indol-3-yl)-2-propenoic acid 5-Methoxyindoleacetate/ Indolelactic acid/ Methyl 1-methoxy-1H-indole-3-carboxylate/ Methyl oxindole-3-acetate	1.22±0.36 1.22±0.36	0.55 0.57	1.10 1.09	-0.30 -0.30
<i>Prenol lipids</i>	Indole-3-ethanol/ Tryptophol Hovenidulcigenin A	1.25±0.19 1.24±0.38	20.91 ns	0.18 -18.55	0.09 0.06
<i>Pyrrolizines</i>	Hydroxysintaxanthin 5,6-epoxide 2,3-Dihydro-1H-pyrrolizine-5-carboxaldehyde	1.25±0.34 1.22±0.32	ns 17.96	-1.32 1.23	-0.006 -0.31

	1-(2,3-Dihydro-1H-pyrrolizin-5-yl)- 2-propen-1-one	1.25±0.19	20.92	0.18	0.09
<i>Quinolines and derivatives</i>	Edulitine	1.22±0.36	0.57	1.08	-0.30
	Graveoline/ Graveoline	1.24±0.40	ns	1.32	0.19
	6-Methylquinoline	1.25±0.19	20.91	0.18	0.09
<i>Steroids and derivatives</i>	Taurochenodesoxycholic acid	1.22±0.45	ns	18.64	0.26
	Withaphysacarpin/14alpha- Hydroxyxocarpanolide/2,3- Dihydrowithanolide E/ Perulactone B	1.22±0.23	ns	-0.28	-17.4
	Lithocholic acid glycine conjugate	1.24±0.37	21.31	ns	0.19
	Cucurbitacide E	1.24±0.38	ns	-1.32	-0.07
	2-Hydroxyestrone sulfate	1.24±0.32	-18.01	-0.36	0.04
	3b-Hydroxy-5-cholenoic acid	1.24±0.35	ns	-1.32	-0.04
<i>Tetrahydrofurans</i>	Tetrahydrofurfuryl acetate/ Botryodiplodin	1.23±0.19	ns	0.17	20.27
<i>Other compounds</i>	2-Aminoacetophenone	1.22±0.32	17.96	1.23	-0.31
	L-Menthone 1,2-glycerol ketal	1.24±0.39	-16.87	ns	0.19
	Avenalumin II	1.24±0.40	ns	1.32	0.19
	Canavaninosuccinate	1.23±0.28	0.21	0.65	19.61
	4-hydroxysphinganine	1.24±0.39	ns	1.32	0.13
	4-Hydroxycyclohexylcarboxylic acid	1.22±0.19	ns	0.17	20.27
	Dihydro-2,4-dimethyl-6-(2- methylpropyl)-4H-1,3,5-dithiazine	1.23±0.33	9.27	9.30	-0.33
	Ethyl levulinate	1.22±0.19	ns	0.17	20.27
	AS 1-5	1.01±0.53	19.91	ns	ns
	N-(2,5- Dihydroxyphenyl)pyridinium(1+)	1.22±0.35	0.52	1.11	-0.31
	(S)-Pterosin K	1.22±0.29	0.20	0.63	-10.32
	6-Chloro-N-(1-methylethyl)-1,3,5- triazine-2,4-diamine	1.22±0.36	0.56	1.08	-0.30

Dictyoquinazol C	1.24±0.32	ns	-1.32	0.03
3-[(5-Methyl-2-furanyl)methyl]-1H-pyrrole	1.25±0.19	20.91	0.18	0.09
(R)-Boschniakine	1.25±0.19	20.91	0.18	0.09
Acetyl-methylpyridine derivatives	1.22±0.32	17.96	1.23	-0.32

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Table 1. Proximate analysis and minerals of edible garlic and elephant garlic cloves. Data were compared with Student *t*-test ($P \leq 0.05$).Significance ns: not significant; *: $P \leq 0.05$; **: $P \leq 0.01$; ***: $P \leq 0.001$ for the interaction of factors.

Proximate composition		Unit	Garlic (<i>Allium sativum</i>)	Elephant garlic (<i>Allium ampeloprasum</i> var. <i>holmense</i>)	Significance
Protein		g 100 g ⁻¹ FW	0.98±0.05	1.22 ± 0,26	*
Carbohydrate	Glucose	g 100 g ⁻¹ DW	3.55±0.52	3.36 ± 0.14	ns
	Sucrose		0.11±0.04	0.11 ± 0.03	ns
Dietary fibre	Neutral detergent fibre (NDF)		9.98±0.16	8.46 ± 0.30	**
	Acid detergent fibre (ADF)		9.09±0.40	8.58 ± 1.09	ns
	Acid detergent lignin (ADL)	%	3.91±0.34	3.25 ± 0.31	ns
	Hemicellulose		0.48±0.33	0.55 ± 0.12	ns
	Cellulose		2.06 ± 0.23	1.29 ± 0.04	**
Minerals	Nitrogen (N)		2.64 ± 0.26	1.88 ± 0.06	**
	Carbon (C)		41.29 ± 0.26	42.38 ± 0.28	**
	Hydrogen (H)		6.56 ± 0.02	6.68 ± 0.04	*
	Sulphur (S)	%	0.38 ± 0.01	0.65 ± 0.18	*
	Phosphorus (P)		1.17 ± 0.04	0.59 ± 0.01	***
	Potassium (K)		1.04 ± 0.11	0.76 ± 0.02	*
	Calcium (Ca)		0.21 ± 0.01	0.23 ± 0.02	ns
	Magnesium (Mg)		0.05 ± <0.01	0.06 ± <0.01	ns
Moisture content		%	62.13 ± 0.57	67.23 ± 0.67	***
Dry matter		%	37.87 ± 0.57	32.77 ± 0.67	***

1 **Table 2.** Bioactive compounds identified by VIP (Variable Importance in Projection) selection method following OPLS-DA model, in the raw
 2 matrix of elephant garlic and common garlic and during the *in vitro* gastrointestinal digestion of the two analysed bulbs. Compounds are provided
 3 together with VIP scores (measure of variable's importance in the OPLS-DA model) and Log Fold-Change in raw material and during different
 4 phases (gastric and pancreatic phases) of the *in vitro* digestion. ¹ns: not significative.

Class	Metabolites	VIP score (OPLS-DA)	LogFC [EG vs CG] raw	LogFC [EG vs CG] gastric	LogFC [EG vs CG] pancreatic
<i>Polyphenols</i>	4-Hydroxybenzoic acid 4-O- glucoside	1.02±0.56	11.42	2.16	9.95
	Rosmarinic acid	1.06±0.25	-18.70	-0.83	ns
	Sinapoylspermine	1.22±0.23	¹ ns	-0.37	-17.73
	Bisdemethoxycurcumin	1.22±0.27	0.35	0.64	2.31
	5-Nonadecylresorcinol	1.00±0.53	21.89	ns	ns
	Phloretin	1.06±0.14	0.49	9.80	ns
	Butein/Naringenin	1.24±0.32	ns	11.23	0.01
	<i>Aminoacids and derivatives</i>	L-Proline	1.02±0.58	-2.56	-1.01
N-acetyl lysine methyl ester		1.23±0.33	9.24	9.28	-0.33
Tryptophan		1.23±0.35	19.61	-0.46	-0.29
Cinnamoylglycine		1.22±0.36	0.57	1.08	-0.30
<i>Benzenoids</i>	(2S,4S)-Monatin	1.22±0.33	0.10	0.64	19.61
	N-Acetylarylamine/ N- benzylformamide/2- Phenylacetamide	1.22±0.32	17.95	1.23	-0.31
	2-(2-Methylpropoxy)naphthalene	1.23±0.33	ns	9.28	-0.33
	4-[(2-Hydroxy-1- naphthalenyl)azo]benzenesulfonic acid	1.23±0.11	0.56	-9.28	ns
	1,1'-[1,12-	1.24±0.35	ns	16.89	0.03

<i>Sulphur containing compounds</i>	Dodecanediylbis(oxy)]bisbenzene N-gamma-glutamyl-S-allylcysteine/ N-gamma-glutamyl-S-trans-(1-propenyl)cysteine	1.01±0.38	-7.01	0.64	2.31
<i>Fatty acyls</i>	Methyl (R)-3-methyl-2-oxopentanoate Pantoyllactone glucoside Sativic acid/ Pinellic acid/9,12,13-TriHOME/(9S,10E,12S,13S)-9,12,13-Trihydroxy-10-octadecenoic acid/5,8,12-Trihydroxy-9-octadecenoic acid/9,10,13-TriHOME	1.22±0.19	ns	0.17	20.27
	Cervonoyl ethanolamide	1.23±0.27	0.23	0.57	18.85
	LysoPC(18:1(11Z))	1.23±0.13	ns	-0.01	-19.08
<i>Glycerophospholipids</i>	LysoPC(20:3(8Z,11Z,14Z)) LysoPC(20:4(8Z,11Z,14Z,17Z)) 1-16:0-2-18:1-phosphatidylcholine 1-18:3-2-18:1-phosphatidylcholine PE-NMe2(16:0/16:0)	1.24±0.38	-7.76	7.80	0.16
	5-(2-Furanyl)-3,4-dihydro-2H-pyrrole	1.24±0.36	-0.65	-18.36	-0.15
	3,4-Dihydro-4-[(5-methyl-2-furanyl)methylene]-2H-pyrrole	1.24±0.39	ns	-1.32	0.15
	3-(1H-Indol-3-yl)-2-propenoic acid	1.24±0.37	ns	-1.32	-0.01
	5-Methoxyindoleacetate/ Indolelactic acid/ Methyl 1-methoxy-1H-indole-3-carboxylate/ Methyl oxindole-3-acetate	1.25±0.16	0.41	17.52	ns
<i>Heteroaromatic compounds</i>	Indole-3-ethanol/ Tryptophol	1.25±0.18	0.34	18.55	0.16
	Hovenidulcigenin A	1.25±0.32	0.15	-1.01	ns
	Hydroxysintaxanthin 5,6-epoxide	1.22±0.32	17.96	1.23	-0.31
<i>Indoles and derivatives</i>	2,3-Dihydro-1H-pyrrolizine-5-carboxaldehyde	1.25±0.19	20.92	0.18	0.09
		1.22±0.36	0.55	1.10	-0.30
		1.22±0.36	0.57	1.09	-0.30
<i>Prenol lipids</i>		1.25±0.19	20.91	0.18	0.09
		1.24±0.38	ns	-18.55	0.06
<i>Pyrrolizines</i>		1.25±0.34	ns	-1.32	-0.006
		1.22±0.32	17.96	1.23	-0.31

	1-(2,3-Dihydro-1H-pyrrolizin-5-yl)- 2-propen-1-one	1.25±0.19	20.92	0.18	0.09
<i>Quinolines and derivatives</i>	Edulitine	1.22±0.36	0.57	1.08	-0.30
	Graveoline/ Graveoline	1.24±0.40	ns	1.32	0.19
	6-Methylquinoline	1.25±0.19	20.91	0.18	0.09
<i>Steroids and derivatives</i>	Taurochenodesoxycholic acid	1.22±0.45	ns	18.64	0.26
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	Cucurbitacide E	1.24±0.38	ns	-1.32	-0.07
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	3b-Hydroxy-5-cholenoic acid	1.24±0.35	ns	-1.32	-0.04
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	Canavaninosuccinate	1.23±0.28	0.21	0.65	19.61
	4-hydroxysphinganine	1.24±0.39	ns	1.32	0.13
	4-Hydroxycyclohexylcarboxylic acid	1.22±0.19	ns	0.17	20.27
	Dihydro-2,4-dimethyl-6-(2- methylpropyl)-4H-1,3,5-dithiazine	1.23±0.33	9.27	9.30	-0.33
	Ethyl levulinate	1.22±0.19	ns	0.17	20.27
	AS 1-5	1.01±0.53	19.91	ns	ns
	N-(2,5- Dihydroxyphenyl)pyridinium(1+)	1.22±0.35	0.52	1.11	-0.31
	(S)-Pterosin K	1.22±0.29	0.20	0.63	-10.32
	6-Chloro-N-(1-methylethyl)-1,3,5- triazine-2,4-diamine	1.22±0.36	0.56	1.08	-0.30

Dictyoquinazol C	1.24±0.32	ns	-1.32	0.03
3-[(5-Methyl-2-furanyl)methyl]-1H-pyrrole	1.25±0.19	20.91	0.18	0.09
(R)-Boschniakine	1.25±0.19	20.91	0.18	0.09
Acetyl-methylpyridine derivatives	1.22±0.32	17.96	1.23	-0.32

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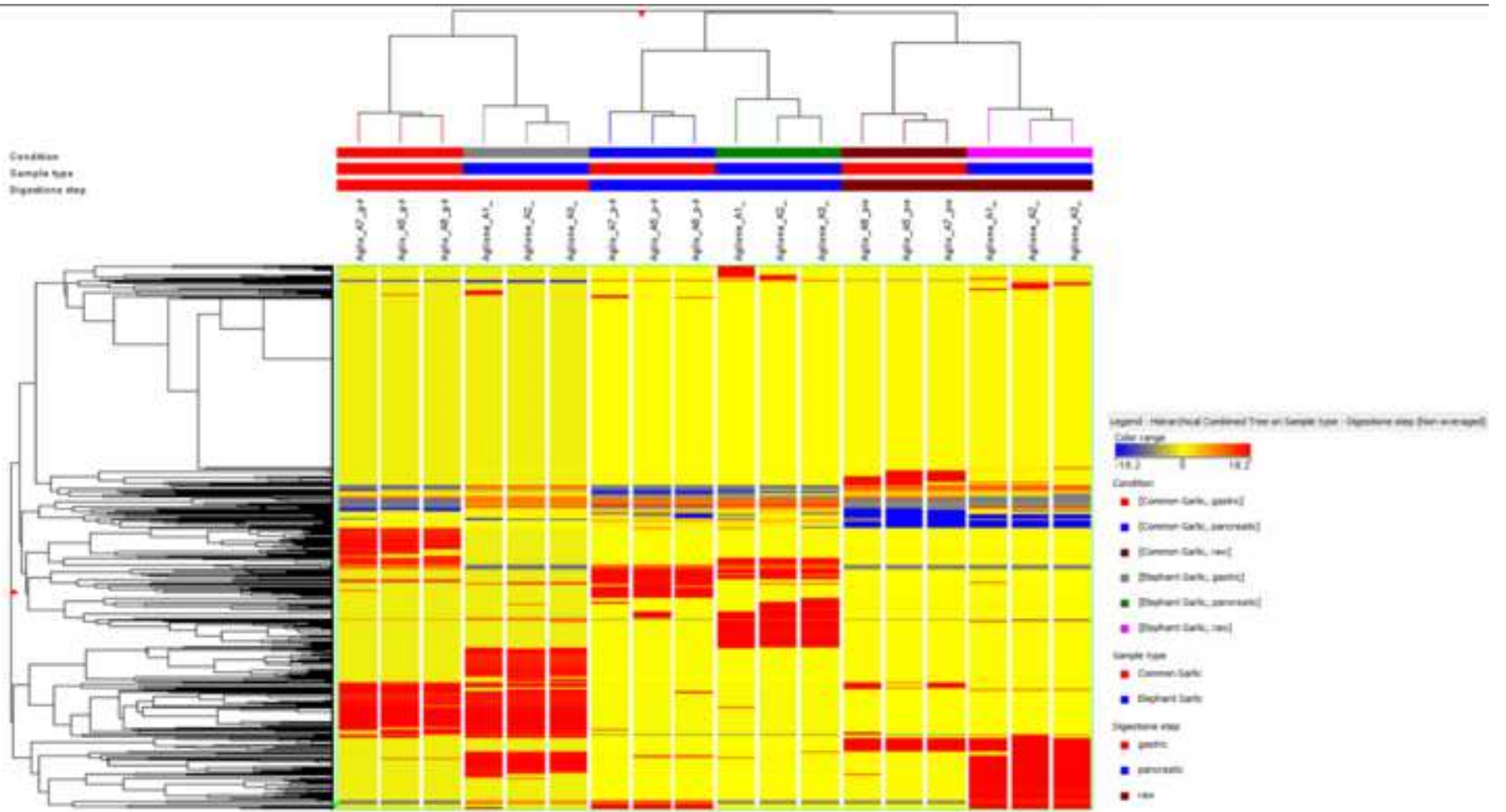
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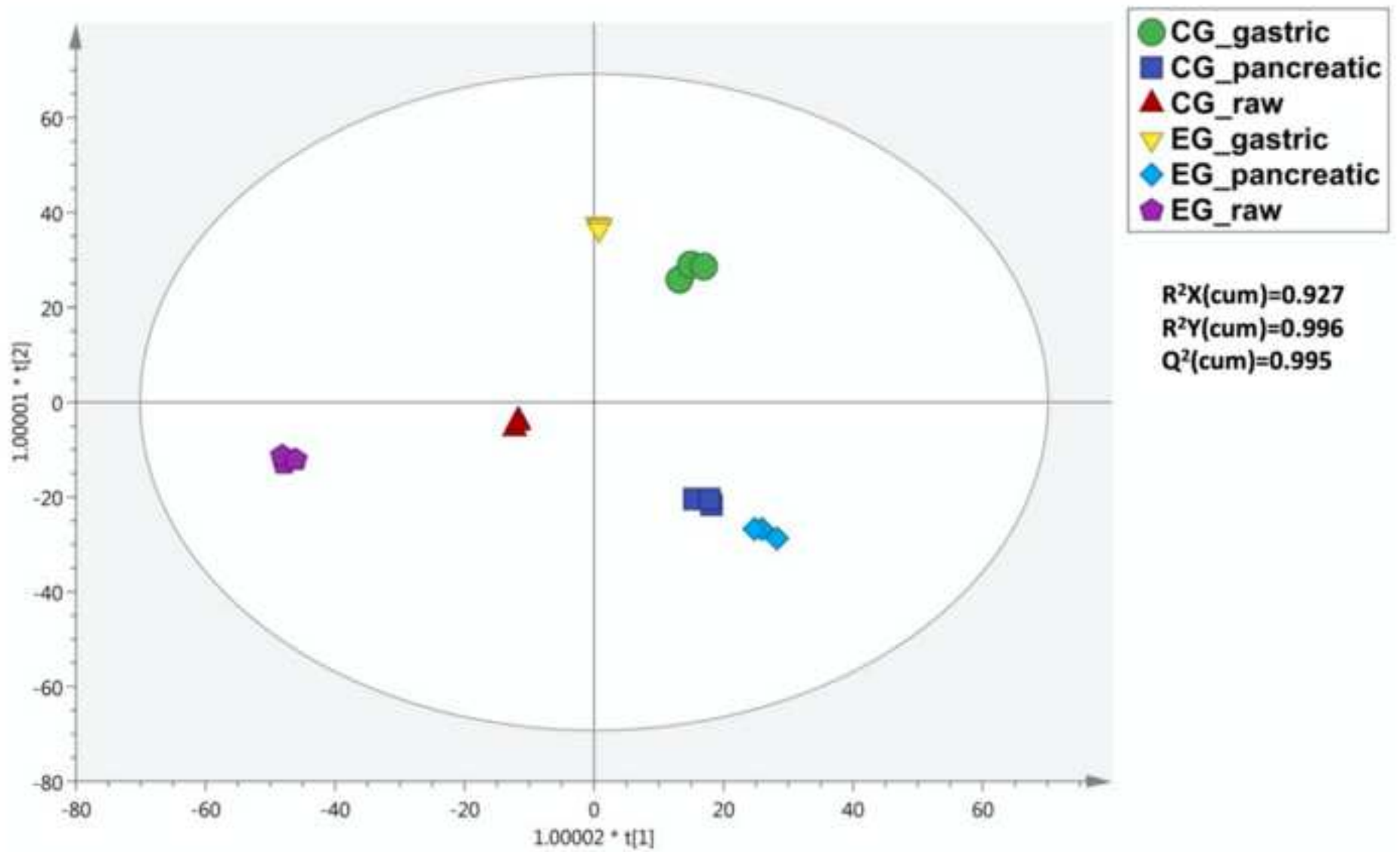


Table S1. Classification of 2745 mass features according to the database FoodDB together with in

Compound	CG gastric	CG gastric
CDP-DG(18:1(11Z)/20:4(5Z,8Z,11Z,14Z))	1	1
CDP-DG(18:1(9Z)/20:4(8Z,11Z,14Z,17Z))	1	1
CDP-DG(18:1(11Z)/20:4(8Z,11Z,14Z,17Z))	1	1
CDP-DG(18:1(9Z)/20:4(5Z,8Z,11Z,14Z))	1	1
Neurine	1.37E+07	1.32E+07
Choline	1.40E+07	1.35E+07
Angiotensin II	1	1
Trigofenoside E1	1	1
Parillin	1	1
Tuberoside B (<i>Allium tuberosum</i>)	1	1
Asparasaponin I	1	1
Protodioscin	1	1
Tuberoside F	1	1
1-Desulfoyessotoxin	1	1
Deltoside	1	1
Avenacoside A	1	1
26-Desglucoavenacoside B	1	1
Dulcin	1	1
Tomatoside A	1	1
Asparagoside G	1	1
Trigofenoside D	1	1
Trigofenoside F	1	1
Tuberoside C (<i>Allium tuberosum</i>)	1	1
Balanitoside	1	1
Tuberoside L	1	1
Matesaponin 2	1	1
Helianthoside A	1	1
Medicoside I	1	1
Pitheduloside K	1	1
Pacific Ciguatoxin 4A	1	1
Cyclolinopeptide D	1	1
Trigofenoside C	1	1
Melongoside P	1	1
Soyasaponin bg	1	1
Bradykinin hydroxyproline	1	1
Eruboside B	1	1
Isoeruboside B	1	1
Cistocardin	1	1
Bradykinin	1	1
Yayoisaponin C	1	1
Tragopogonsaponin Q	1	1
Camellidin II	1	1
Capsianoside III	1	1

Table S2. Volcano plot comparing raw elephant garlic vs common garlic by coupling ANOVA (**Moderated T-Test [Elephant Garlic, raw] vs [Common Garlic, raw] $P \leq 0.01$ $FC \geq 5.0$** **Compound**

L-Proline
5-Acetyl-2,3-dihydro-1,4-thiazine
L-2-Amino-3-(1-pyrazolyl)propanoic acid
trans-Carvyl acetate
Rhubafuran
(R)C(S)-Alliin
Polyvidone
6-Acetylfuranofukinol
Urodiolenone
Citflavanone
Cyclocalopin F
Sterebin A
(1R*,3S*,3'R*)-1,2,3,4-Tetrahydro-1-(2-thio-3-pyrrolidinyl)-beta-carboline-3-carboxylic acid
L-Menthone 1,2-glycerol ketal
N-gamma-Glutamyl-S-allylcysteine
N-gamma-Glutamyl-S-cis-(1-propenyl)cysteine
Rosmarinic acid
Isotriglochinin
Irigenin
Gerberinol
2-Hydroxyestrone sulfate
Artonin K
4'-phosphopantetheine
Gonyautoxin V
Gingerenone B
Dihydrofukinolide
Myristicanol B
Boviquinone 4
2-(2-Methylbutanoyl)-9-(3-methyl-2E-pentenoyl)-2b,9a-dihydroxy-4Z,10(14)-oplopadien-3-or
Clausarinol
Lusitanicoside
Erinacine D
ent-Epiafzelechin-(2alpha-7,4alpha-8)-catechin
DG(18:3(9Z,12Z,15Z)/18:0/0:0)
Ramontoside
Epiafzelechin-(4alpha-8)-pelargonidin 3'-glucoside

Compound

N-Methylcalystegine C1
Xylopine
Alkaloid A6
Cyclolinopeptide D
Bradykinin hydroxyproline

Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: