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6 Title: Effect of natural antioxidants from grape seed and chestnut in combination with
7 hydroxytyrosol, as sodium nitrite substitutes in Cinta Senese dry-fermented sausages

# 8 Authors:

9 Chiara Aquilani<sup>a\*</sup>, Francesco Sirtori<sup>a</sup>, Monica Flores<sup>b</sup>, Riccardo Bozzi<sup>a</sup>, Benedicte Lebret<sup>c</sup>, Carolina
10 Pugliese<sup>a</sup>

<sup>a</sup>DISPAA - Department of Agrifood Production and Environmental Sciences, Animal Science
 Section, Universita' degli Studi di Firenze, 50144 Firenze, Italy

<sup>13</sup> <sup>b</sup>IATA - Institute of Agrochemistry and Food Technology (IATA-CSIC), Paterna, Valencia, Spain

14 °PEGASE, INRA, Agrocampus-Ouest, 35042 Rennes, France

# 15 **\* Corresponding author:** <u>chiara.aquilani@unifi.it</u>

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## 17 Abstract:

Dry-fermented pork sausages, from Cinta Senese local breed, were manufactured replacing sodium nitrite (NIT) with two mixtures of natural antioxidants consisting of: i) grape seed extract and olive pomace hydroxytyrosol (GSE); ii) chestnut extract and olive pomace hydroxytyrosol (CHE). The effects on physical-chemical, aromatic and sensory traits, as well as the microbiological safety, were tested. Nitrite replacement lowered the pH in GSE and CHE samples and resulted in several differences in physical traits between CHE and NIT samples. *Listeria monocytogenes, Salmonella*  and *Clostridium botulinum* were not found in any samples. GSE and CHE mixtures showed a slightly lower antioxidant activity. Volatile profile showed a similar aromatic profile among the three treatments with differences mainly to abundance of the single compounds, indicating that replacement of nitrite by natural antioxidants did not affect the overall aroma profile, as outlined by olfactometry results. In addition, the replacement did not affect the overall acceptability, except for color-related traits, underscored in GSE and CHE products.

Keywords: Volatile compounds; meat quality; GC-olfactometry; local pig breed; lipid oxidation;
pork products

# 32 **1. Introduction**

Dry cured meat products are typical of the Mediterranean area and they represent a high-value 33 production in European countries, considering that curing process allows extension of meat shelf-34 life (Marco et al., 2006) and leads to typical pork products with specific eating quality and regional 35 identity (Pugliese and Sirtori, 2012). In Southern Europe, salami and dry-fermented sausages, are 36 37 generally characterized by slowly air-drying and mold-ripening (Flores, 1997). This curing process leads to peculiar characteristics and flavors that are widely appreciated by consumers; but is also 38 related to longer curing times that may cause higher lipid oxidation levels. Moreover, natural 39 40 fermentation, avoiding the addition of lactic acid-producing starter cultures, is more susceptible to the growth of harmful bacteria, such as Listeria monocytogenes or Clostridium botulinum (Lücke, 41 2000). Thus, to avoid a severe deterioration of nutritive and organoleptic attributes, as well as to 42 ensure food safety, several synthetic food preservatives are commonly included. Among them, the 43 most used are nitrites and nitrates (Hammes, 2012). Nitrite positively affects color, inhibits the 44 45 growth of pathogenic bacteria, contributes to the development of typical cured meat flavor and delays oxidative rancidity (Marco et al., 2006). Despite their effectiveness as curing agents, the 46 nitrite/nitrate intake represents a risk to human health, i.e. the formation of carcinogenic 47 nitrosamines is one of the most current concerns (De Mey et al., 2017). Several studies have 48

focused on nitrate/nitrite reduction or substitution (Purriños et al., 2013; Özvural and Vural, 2014; 49 50 Pateiro et al., 2015), but the main issue remains finding an alternative able to address the multiple activities they perform. Up until now, most of the alternatives proposed are plant extracts, largely 51 obtained from agricultural by-products. These compounds are very rich in polyphenols, flavonoids 52 and terpenoids and are able to perform a double antioxidant-antimicrobial functions (Falowo et al., 53 2014; Hygreeva et al., 2014; Shah et al., 2014). These compounds might also constitute a great 54 55 opportunity to exploit agricultural by-products, which otherwise would be wasted. The aim of this study was to assess the feasibility of producing dry-fermented sausages by replacing sodium nitrite 56 with natural antioxidants while trying to maintain quality traits. Grape seed extract, chestnut extract 57 58 and hydroxytyrosol (extracted from defatted olive pomace), were chosen due to their great availability as by-products of important Tuscan agricultural products. Moreover, among the 59 investigated plant extracts, they have shown an interesting potential both for antioxidant activity 60 61 and microbial inhibition. This innovation also aimed to valorize Cinta Senese, a local pig breed strongly linked to the Tuscan region. 62

63 **2. Materials and methods** 

# 64 **2.1. Antioxidant mixtures**

The natural antioxidants employed in the present studies were provided by Phytolab (Sesto Fiorentino, Florence, Italy). They consisted of grape seed and chestnut extracts, tocopherol and hydroxytyrosol (extracted by defatted olive pomace). The manufacturer provided the phenolic profile (Table 1), total phenolic content and antiradical scavenging activity ( $EC_{50}$ ) (Table 2) of each extract. The grape seed and chestnut extracts were combined with the same amount of hydroxytyrosol and tocopherol to form two different mixtures; grape seed (GSE) and chestnut (CHE) mixtures.

# 72 **2.2. Sausages manufacturing**

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In an industrial plant (Azienda Agricola Savigni, Pistoia, Italy), 24kg of pork lean and 6kg of 73 74 subcutaneous backfat from Cinta Senese pig breed were minced and equally divided in three batches. Salt (23g/kg), sucrose (35g/kg) and black pepper (0.2g/kg) were added to each batch 75 following the recipe traditionally used by the manufacturer. Thirty ppm of sodium nitrite (E250) 76 were added to the first batch to constitute the control (NIT). In second batch, 10g/kg of GSE 77 mixture were used to replace sodium nitrite, while 10g/kg of CHE were added to the third batch. 78 79 Sausages were weighed, dried at 28°C and RH 85% for 4 days and then ripened 21 days (T 13°C, RH 70%). Once ripened, six samples of each batch were collected, pH, color, and processing loss 80 were immediately measured. Samples were vacuum packed and stored at -80°C for physical, 81 82 chemical and aromatic analysis. Another 3 samples of each batch were stored at 4°C to be employed for sensory analysis the following day. This design was replicated to have two totally 83 independent batches for each treatment. 84

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## 2.3. Physical, chemical and microbiological parameters

At the end of ripening, physical parameters were assessed on 12 samples of each batch (6 for each 86 replication). Sausage pH was measured at room temperature (20 °C) using a pH meter Crison 87 GLP21 (Barcelona, Spain), the instrument was introduced in a sausage portion. Color (L\*, a\* and 88 b\*) was determined by a Minolta Chromameter CR-200 (Tokyo, Japan) immediately after slicing. 89 aw was measured following the method ISO 21807:2004. Two 10 mm-thick and 10 mm-width slices 90 of each sample, were cut and immediately analyzed at room temperature (22 °C), using a Zwick 91 Roell Z2.5 apparatus (Ulm, Germany) with a loading cell of 1 kN at the crosshead speed of 1 92 93 mm/sec. Texture profile analysis (TPA) was performed assessing the following parameters: hardness, cohesiveness, gumminess, springiness and chewiness. Moisture was determined by 94 95 lyophilizing to constant weight 40g of sample, according to AOAC methods (1990). Weight loss was measured as the difference between weight at time zero and end of ripening (after 24 days). 96 Total protein, fat and ash contents were determined following AOAC (1990) methods. Lipid 97

oxidation was determined according Vyncke (1970), using a PerkinElmer Lambda EZ150 98 99 spectrophotometer (Waltham, MA, USA). Results were expressed as mg of malondialdehyde (MDA)/kg of samples. Fatty acids were determined using a Varian GC-430 apparatus equipped 100 with a flame ionization detector (FID) (Palo Alto, CA, USA) as reported by Sirtori et al. (2015). 101 The individual methyl esters were identified by their retention time using an analytical standard 102 (F.A.M.E. Mix, C8-C22 Supelco 18920-1AMP). Response factors based on the internal standard 103 (C19:0) were used for quantification and results were expressed as mg/100g of sample. The fatty 104 acid content was reported as saturated (SFA), monounsaturated (MUFA) and polyunsaturated 105 (PUFA) fatty acids. Microbiological analyses were carried out in an external accredited laboratory 106 107 to determine the products' safety. The following bacteria were investigated: Escherichia coli (ISO 108 16649-2:2001), Listeria monocytogenes (UNI EN ISO 11290-1:2005), coagulase positive Staphylococcus spp. (UNI EN ISO 6888-1:2004), Clostridium botulinum (ISO 15213:2003) and 109 Salmonella spp. (UNI EN ISO 6579:2008). 110

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## 2.4. Volatile compounds analysis

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## 2.4.1. Gas chromatography-mass spectrometry analysis (GC-MS)

Solid-phase microextraction (SPME) and GC-MS analysis were performed following the method 113 114 described by Corral, Salvador, & Flores (2013) using a 85 µm Carboxen/Polydimethylsiloxane (CAR/PDMS) fiber (Supelco, Bellefonte, PA) installed in a Gerstel MPS2 multipurpose sampler 115 (Gerstel, Germany) and an Agilent HP 7890 series II GC with an HP 5975C mass selective detector 116 117 (Hewlett-Packard Palo Alto, CA, USA). The volatile compounds (VOCs) detected were identified by comparison with mass-spectra from the library database (Nist'05), linear retention index (van 118 Den Dool and Dec. Kratz, 1963) and by comparison with authentic standards. The quantification of 119 volatile compounds was done in SCAN mode using either total or extracted ion chromatogram (TIC 120 or EIC) on an arbitrary scale. 121

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## 2.4.2. Gas chromatrography-olfactometry analysis (GC-O)

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A gas chromatograph (Agilent 6890, USA) equipped with an FID detector and sniffing port (ODP3, 123 124 Gerstel, Mülheim an der Ruhr, Germany) was used to analyze aroma compounds extracted by SPME as described by Corral et al. (2013). The detection frequency (DF) method was used to 125 estimate the aromatic impact of each volatile and each assessment was carried out according 126 to Olivares, Navarro, & Flores (2011). Four trained panelists evaluated the odors from the GC-127 effluent. Each assessor evaluated 3 sausages for a total of 12 assessments, the final DF was obtained 128 129 by summing the 12 sniffings. The detection of an odor by less than three assessors was considered noise. Compounds were identified by comparison with mass spectra, with linear retention indices 130 of authentic standards injected in the GC-MS and GC-O, and by coincidence of the assessors' 131 132 descriptors with those reported by Burdock (2010).

## 133 **2.5. Sensory analysis**

Sensory analysis was carried out in an equipped laboratory by 8 trained panelists using a 134 quantitative-descriptive analysis method. Fourteen attributes (grease appearance, abnormal colors, 135 136 firmness, color uniformity, redness, cured meat flavor, off odor, salty, rancid, off flavor, hardiness, juiciness, aftertaste, general acceptability) were evaluated, each attribute was scored in a 10 cm 137 non-structured line (Pugliese et al., 2010). Select subjects underwent an introductory session, where 138 the testing procedures and the chosen sensory traits were discussed using two types of comparable 139 commercial products. During three sessions, panelists evaluated a total of 9 sausages (3 samples x 3 140 treatments) identified by an alphanumerical code. The sausages were divided in 0.5cm-thick x 2cm-141 diameter slices and two slices of each samples were randomly served to judges at room temperature 142 (20°C). Panelists were invited to eat a cracker and drink a glass of water between samples. 143

## 144 **2.6. Statistical analysis**

145 Data were analyzed by SAS software. Two-way ANOVAs were performed on physical and146 chemical data according to the following model:

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147  $Y_{ijk} = \mu + T_i + B_j + \varepsilon_{ijk}$ 

148 Where  $\mu$  is the mean, T is the i<sup>th</sup> treatment, B is the j<sup>th</sup> batch and  $\varepsilon$  is the error. For sensory data, 149 effect of panelist was included in the previous model. The interaction between Treatment and Batch 150 factors was tested but being not significant, it was not included in the model.

Volatile compounds data were also analyzed by a multivariate approach to determine the presence 151 of characteristic compounds able to be allocated to samples among different treatments. A stepwise 152 discriminant analysis (SDA) was first used to reduce the space-variables, selecting the subset of 153 variables that better discriminated groups. Canonical discriminant analysis (CDA) were performed 154 155 using SDA selected variables, resulting in 2 new variables, called canonical functions (CAN1, CAN2). They consisted of a series of canonical coefficients (CC) that indicate the partial 156 contribution of each variable in composing the CANs. The greater the CC, the more the variable 157 contributes to CAN composition. 158

## 159 **3. Results and discussion**

## 160 *3.1. Physical, chemical and microbiological parameters*

The major foodborne pathogens (Escherichia coli, Listeria monocytogenes, Staphilococcus spp., 161 Clostridium spp and Salmonella spp.) were absent or below the limit required (Reg CE/2073/05) in 162 163 all samples (Table 3). Several studies on plant phenolic suggest these components have antimicrobial activity (Kao et al., 2010; Mujić et al., 2014; Fasolato et al., 2016). Further studies 164 are, however, required to determine the effectiveness of the studied antioxidants against the 165 development of the main foodborne pathogens. The a<sub>w</sub> values (Table 4) recorded for GSE and CHE, 166 being below 0.89, contributed to control pathogenic organisms development (Toldrá and Flores, 167 168 2014). Moisture, fat, protein and ash contents and weight loss were not affected by treatment and they were in line with values reported for dry-cured sausages (Olivares, Navarro, & Flores, 2015; 169 Ribas-Agustí et al., 2014; Škrlep, Čandek-Potokar, Tomažin, Batorek Lukač, & Flores, 2017), 170 171 except for weight loss, which was slightly greater in the present study; likely the smaller diameter

of samples could have enhanced the water loss during ripening. The pH observed was in the range 172 173 reported for natural fermented meat products at a similar curing time (Özvural and Vural, 2014; Škrlep et al., 2017); indeed, they were characterized by a higher pH compared to commercial 174 products (Hospital et al., 2015; Montanari et al., 2018). Moreover, significant differences among 175 treatments were found for pH, with lower values observed for GSE and CHE samples, suggesting 176 that Lactobacillus (LAB) growth, that takes place during the first fermentation phase, could be 177 178 slightly promoted in these products. Indeed, low or nitrites-free sausages showed an increased presence of LAB (Hospital et al., 2015). Growth of LAB was not, however, assessed in the present 179 experiment, and further studies will be required to assess effects of GSE and CHE. Concerning 180 181 color attributes, L\* was not affected by treatment, a\* showed significant greater values in GSE and NIT samples than in CHE ones, while b\* was significant higher in NIT compared to the modified 182 products. A change in a\* was expected considering the role nitrites play in nitrosomyoglobin 183 184 formation, the characteristic red curing pigment (Hammes, 2012). Since neither chemical composition nor oxidation resulted in significant differences among groups, a different pathway for 185 red color formation in GSE samples should be considered. A stable red color compound called Zn-186 protoporphyrin, derived from the substitution of heme iron with zinc, has been observed in Parma 187 ham, an Italian nitrite/nitrate-free ham (Wakamatsu et al., 2004a). Up to now, mechanisms leading 188 189 to its formation are not well-known (Hammes, 2012), but the absence of nitrites, low levels of oxygen, meat endogenous enzymes as well as microorganism, are all factors that may contribute to 190 its formation (Wakamatsu et al., 2004b). Apart from the absence of nitrites, some compounds 191 192 contained in grape seed extract may have promoted the Zn-protoporphyrin formation for the GSE group, while no formation occurred in the CHE samples. These results are partially in agreement 193 with those reported by Lorenzo, González-Rodríguez, Sánchez, Amado, & Franco (2013), althought 194 their study was conducted on chorizo where pigmentation due to added paprika may have 195 interfered. 196

According to TPA results (Table 5), cohesiveness, springiness and chewiness were affected by 197 198 nitrite replacement, being highest in CHE samples, lowest in NIT, whereas GSE samples were similar to both. Since no differences in moisture and weight loss were found among the treatments, 199 the results obtained were attributable basically to the differences in pH, which, declining, causes the 200 aggregation of myofibrillar proteins and leads to gel formation (Lücke, 2000). The higher pH of 201 202 NIT samples likely inhibited this phenomenon thus reducing the sausage cohesiveness and 203 chewiness. The results reported are partially in agreement with Lorenzo et al., (2013), who also noticed the highest chewiness values for chorizo with added chestnut extract and ripened for 19 204 205 days, compared to the same product manufactured with GSE or synthetic antioxidant (BHT in this 206 case).

The groups did not differ in SFA, MUFA and PUFA contents. Their relative amounts reflect those 207 of fresh pork composition, slightly richer in MUFA than SFA, with PUFA being approximately a 208 third of either SFA or MUFA categories (Škrlep et al., 2017). PUFA can also be considered an 209 210 indicator of meat oxidative status, due to their double bonds being preferred substrates for oxidative 211 reactions (Pateiro et al., 2015). Results suggest that the natural antioxidants employed were as 212 effective as nitrites in control lipid oxidation during manufacturing and ripening. This is supported by TBARs results, showing no significant differences among treatments, however, further studies 213 214 will be required to evaluate the antioxidant activity during the shelf-life. Nitrites exert their antioxidant activity in cured meat by forming the myoglobin-stable compounds and making the iron 215 inaccessible for oxidation (Riazi et al., 2018); phenolic compounds instead, follow different 216 pathways, acting as hydrogen donors. The phenolic hydroxyl groups intercept the free radicals to 217 form stable end-products, interrupting and avoiding further lipid oxidation, especially of 218 219 unsaturated FAs (Jayaprakasha et al., 2003). To the best of our knowledge, few data are available about dry-fermented pork sausages with added natural extracts, but the great variability of these 220 traditional products makes comparisons difficult. The efficacy of hydroxytyrosol in preventing lipid 221

oxidation was reported by Cofrades et al., (2011) for n-3 enriched frankfurters, while Lorenzo et al.,
(2013) observed comparable TBARS values in Spanish chorizo with added BHT, grape seed extract
or chestnut extract.

# 225 *3.2. Volatile profile and olfactometry*

Ninety-one VOCs were identified by HS-SPME-GS-MS (Table 6). The most abundant groups originated from spices (51-61%) and carbohydrate fermentation (30-39%), followed by amino acid degradation (6-7%), while VOCs derived from lipid  $\beta$ -oxidation and lipid oxidation processes represented 1% of total extracted areas.

230 Among the 14 VOCs related to lipid auto-oxidation, 5 showed significant differences with NIT resulting in the lowest, while GSE and CHE products showed intermediate or higher abundances. 231 These VOCs originate by autocatalytic fat oxidation and involves mostly unsaturated fatty acids. 232 Among the identified VOCs, hexanal is of key-importance to better outline the products' oxidation 233 status. The correlation between this compound and lipid oxidation is well-known and its low 234 235 perception threshold makes hexanal an important contributor to overall aroma (Marco et al., 2006). The higher hexanal content in GSE and CHE samples than NIT is consistant with TBARs results, 236 suggesting greater PUFA oxidation, even if the extent was limited and did not affect the parameters 237 238 previously examined. The total lipid auto-oxidation values confirmed these differences, with NIT being the lowest, GSE the highest and CHE similar to both, likely related to a higher EC<sub>50</sub> for GSE. 239 Hence, even though the phenolic extracts used contributed to maintain lipid oxidation below the 240 perception threshold of rancid flavor, they appeared less effective than nitrites in controlling lipid 241 oxidation. Partially in agreement with this, Purriños et al. (2013) reported grape seed was a less 242 243 effective antioxidant in chorizo, but on the contrary, chestnut extract was found to have higher antioxidant activity than BHT. 244

Spice derived VOCs were the major group, due to the use of black pepper. Indeed, limonene, a compound particularly abundant in pepper (Moretti et al., 2004) represented approximately half of

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the total amount for each treatment. As the same amount of pepper was added to all treatments, differences among the 3 groups might be due to several external factors such as an irregular distribution of the ground pepper in the raw matrix (Montanari et al., 2018), as well as a heterogeneity of the spices themselves caused by different grinding techniques and/or storage times that could have led to differential oxidative status, odorant losses and sensory attribute changes (Orav et al., 2004; Liu et al., 2013).

253 Microbial enzymes degrade free fatty acids through β-oxidation reactions, generating methyl ketones as final products (Flores & Olivares, 2007). Two VOCs belonging to this group were 254 identified, but significant differences were found only for total abundance, which was highest for 255 256 NIT, the lowest in GSE and CHE samples were similar to both. Since the main microbial populations were not examined in this study, the differences in volatile development due to 257 258 microbial fermentation cannot be explained directly. However, the role of genus staphylococcus in incomplete  $\beta$ -oxidation are well-known, so the presence of 2-pentanone and 3-octanone were likely 259 related to the presence of these bacteria (Chen et al., 2017). 260

Contrarily to Chen et al., (2017) and Marco et al. (2006), only one ester was observed. However, they worked on fermented sausages manufactured with starter cultures containing staphylococci strains. Staphylococci promote the formation of esters with staphylococci esterase activity being one of the main factors leading to ester formation in dry-fermented sausages (Wang et al., 2016).

Among VOCs generated by bacterial metabolism, products of carbohydrate fermentation form an important group, consisting in 13 identified compounds. Acetaldehyde, 2,3-butanedione and 2,3butanediol were higher in GSE samples than in CHE and NIT; ethyl alcohol and butanoic acid were lowest in NIT samples, while 3-hydroxy-2-butanone was lowest in CHE samples. VOCs from carbohydrate metabolism were generally the highest for GSE. The great abundance of 2-butanone is remarkable, considering that this compound is known as a by-product LAB metabolism (Montanari et al., 2018), originating from 2,3-butanedione (significantly higher in GSE). Then, 2-butanone is 272 reduced to 2-butanol and 3-hydroxy-2-butanone, that again, was significantly higher in GSE
273 samples and preferentially formed in small diameter sausages (Montanari et al., 2018).

The last group of VOCs related to bacterial metabolism consisted of 19 VOCs from amino acid 274 275 degradation. 2-methylpropanal and ethylbenzene were higher in CHE samples, while the lowest amounts of 3-methylbutanol and 2-acetyl-1-pyrroline were observed for natural antioxidant 276 products and benzeneacetaldehyde and benzylalcohol were higher in GSE samples than for CHE 277 and NIT. The highest total amino acid degradation products were observed in NIT samples, 278 followed by CHE and then GSE. This was likely due to 3-methylbutanol, whose content almost 279 doubled in NIT samples while toluene, the most abundant compound, was similar for the three 280 groups. The compounds observed are characteristic of dry-fermented sausages, being reported by 281 several authors (Marco et al., 2006; Corral et al., 2013; Škrlep et al., 2017). 282

Despite the variability within each group of VOCs, total amounts of microbial metabolites suggest a greater development of microflora in NIT samples that might be related to the antimicrobial activity of phenolic extracts during ripening. Several studies have reported phenolic compounds diffuse into bacterial cells walls and interact with cytoplasmatic proteins, affecting Gram positive bacteria and, particularly, Gram positive cocci (Jayaprakasha et al., 2003; Fasolato et al., 2016; Riazi et al., 2018). It is worth noting that the main populations involved in sausage fermentation processes, LAB and Staphylococci, are both Gram positive bacteria.

The role each compound plays in defining the aromatic profile strongly depends on its abundance and on its perception threshold (Olivares et al., 2015). During the GC-O sessions, 31 aroma notes were perceived by trained assessors (Table 7). Seven aroma compounds were associated with spices, 1 to lipid beta-oxidation, 4 to carbohydrate fermentation, 1 to esterase activity, 11 to amino acid degradation, 4 to lipid auto-oxidation, 2 to unknown compounds (not identified with any of the VOCs identified by GC-MS) and 1 to contaminants. Considering their DF value as an aroma impact index, 11 VOCs had a DF higher than 8. Despite the differences outlined by SPME-GC-MS

analysis, panelists did not detect any differences in the olfactometric profile of the three groups. 297 298 This is likely because all the identified VOCs were observed in the three groups and the main differences among GSE, CHE and NIT samples were attributable only to differences in the single 299 compounds abundances. As a consequence, GC-O data were displayed combined in a single 300 aromatic profile (Table 7). Most of the identified compounds were previously observed as recurrent 301 in dry-fermented sausages (Flores & Olivares, 2015; Söllner and Schieberle, 2009; Schmidt and 302 303 Berger, 1998). Amino acid degradation compounds have a key-role in flavor development, contributing with malty, fruity, sweaty flavors and ripened aroma (Hospital et al., 2015; Chen et al., 304 305 2017). Indeed, more than half of SPME-GC-MS identified VOCs were also observed in GC-O 306 sessions as odor active compounds in Cinta Senese dry-fermented sausages. They accounted for one 307 third of the compounds forming the GC-O profile and had high DF values. Among them, 2 acetyl-1-pyrroline and methional are considered as the most potent odor active compounds in dry-308 309 fermented sausages (Corral, Leitner, Siegmund, & Flores, 2016; Söllner and Schieberle, 2009). Also 3-methylbutanoic acid is considered a potent aroma contributor, giving cheesy, lactic and fatty 310 notes (Flores & Olivares 2015), while 2,5-dimethylpyrazine is related to meaty and cooked potatoes 311 312 notes.

The second most represented VOC group were spice-derived, especially  $\alpha$ -terpinene,  $\beta$ -myrcene 313 and terpinolene, which were previously reported as odor active compounds (Olivares et al., 2015; 314 Schmidt & Berger, 1998). Another important group was composed of lipid oxidation VOCs, among 315 them hexanal was the most potent odorant (Marco, Navarro, & Flores, 2007; Olivares et al., 2015; 316 Söllner and Schieberle, 2009; Schmidt and Berger, 1998). Indeed, hexanal produces fresh and green 317 notes (Table 7), but it turns to rancid notes as its abundance increases. Lastly, also 2,3-butanedione, 318 319 derived from carbohydrate fermentation, due to its low threshold (about 4 µg/l) (Hospital et al., 2015), was an important aroma contributor, characterized by buttery-sweet notes and a DF of 10. 320

Figure 1 displays the 19 compounds identified by SDA. The selected compounds were able to 321 322 discriminate the three treatments. Can1 accounted for a great part of variance, separating CHE from GSE and NIT, while Can 2 sharply divided GSE and CHE from NIT. Multivariate analysis showed 323 how lipid auto-oxidation compounds, comprising half of the compounds identified by SDA, were 324 central in differentiating the three groups. Tetradecane, ethylbenzene and 3-octanone having the 325 greatest negative Can1 scores, were considered mainly responsible for separating GSE and NIT 326 327 from CHE, in agreement with ANOVA results. Concerning Can2, compounds with higher weighing were decane and octane for GSE and CHE, while 3-methylbutanol seems to characterize NIT 328 samples. Among the 19 compounds identified, 8 were also perceived by GC-O panelists. However, 329 330 considering their DF and CCs together, only hexanal and 2.5-dimethylpyrazine might have an 331 effective potential in discriminating the groups also from a sensory point of view.

# 332 3.3. Sensory analysis

Figure 2 shows sensory results. Abnormal colors, off-flavors, off-odors and rancid were scored as 0 333 334 (not present) and were not shown. As expected, the most affected traits were color related, with 335 both color uniformity (P < 0.01) and redness (P < 0.01) scoring lower for GSE and CHE samples compared to NIT ones, while firmness was was scored highest for CHE samples (P < 0.05), 336 agreeing with TPA results. Likely, the lower moisture and fat contents observed in CHE could have 337 affected the firmness, even if neither moisture nor fat, significantly differed among treatments. 338 Despite the differences in lipid auto-oxidation VOCs, no perceivable rancid notes were detected by 339 panelists, as the level of MDA found in the samples did not exceed the organoleptic perception of 340 341 lipid oxidation (Campo et al., 2006). Effects of adding grape seed and chestnut extracts to dryfermented sausages have not, however, always been positive. Ribas-Agustí et al. (2014) reported 342 343 that panelists discarded grape seed extract added products, as there were judged to be abnormal compared to control samples. Similarly, Özvural and Vural (2011) observed a decrease in overall 344

acceptability of frankfurters with grape seed extract added, even when products manufactured with
 concentrations lower than 0.05% resulted in scores similar to control.

# 347 **4.** Conclusions

The results on VOCs profiles suggested a greater antimicrobial activity of natural antioxidant 348 mixtures (GSE and CHE) compared to sodium nitrite (NIT), likely due to their phenolic 349 constituents; further none of the main foodborne pathogens were found in any sample. No 350 significant differences among treatments were found for lipid oxidation, even if lipid auto-oxidation 351 VOCs suggested a slightly lower antioxidant activity of GSE and CHE compared to sodium nitrite. 352 353 Despite the differences in single VOCs abundances, the replacement did not affect the overall aroma profile, as outlined by GC-O results and sensory analysis. Some differences in instrumental 354 color and texture negatively affected GSE and CHE products, but the overall acceptability was not 355 influenced. GSE and CHE effects on microbiota in dry-fermented sausages should be studied in 356 depth, however, the results so far indicated that tested antioxidants are valid alternatives to sodium 357 nitrite in Cinta Senese dry-fermented sausages. 358

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Olive Pomace (g/L) (hydroxytyrosol)	(e )			Chestnu (mg/g)	t
Hydroxytyrosol	11.65	( <b>mg</b> / Gallic acid	0.01	Vescalin	9.34
Tyrosol & hydroxytyrosol derived compounds	15.13	Catechin B3 (dimers)	2.22	Castalin	8.99
Verbascosid	5.84	Catechin	11.07	Pedunculagin I	3.88
		Catechin (trimers)	3.21	Monogalloil glucose I	3.58
		Catechin B6 (dimers)	2.61	Gallic acid	18.50
		Catechin B2 (dimers)	5.37	Monogalloil glucose II	2.73
		Epicatechin	13.62	Roburin D	10.51
		Catechina trimer	3.71	Vescalagin	32.15
		Epicatechin gallate (PM 730)	6.65	<i>C</i> -glucoside tergallic dehydrate	2.73
		Epicatechin gallate (PM 442)	6.10	Castalagin	31.03
		Oligomers (tetramers)	54.88	Digalloil glucose I	10.03
		Epicatechin gallate (PM 882)	180.65	Digalloil glucose II	2.09
		Epicatechin gallate oligomers (trimers)	382.97	Hydrolyzable tannin <i>m/z</i> 1085	8.05
		Epicatechin gallate oligomers (trimers)	149.66	Trigalloil glucose I	4.61
		、		Trigalloil glucose II	6.74

489 Table 1. Phenolic profile of olive pomace and defatted grape seed and chestnut extracts.

Tetragalloil glucose	2.05
Ellagic acid	4.08

490

491

492 Table 2. Total phenolic content and radical scavenging activity of natural antioxidant constituting493 the mixtures

Total phenolic content	Antiradical scavenging
	activity (EC50)
822.709 (mg/g)	0.147
161.091 (mg/g)	0.085
32.62 (g/l)	0.196
-	0.184
	822.709 (mg/g) 161.091 (mg/g)

Table 3. Microbiological safety parameters on Cinta Sense dry-fermented sausages manufactured
with natural antioxidant (GSE=grape seed extract; CHE=chestnut extract) as replacement of sodium
nitrite (NIT).

	GSE	CHE	NIT
Escherichia coli	<10	<10	<10
Listeria monocytogenes	-	-	-
Coagulase positive	<10	<10	<10
Staphilococcus spp.	<10	<10	<10

Sulfite-reducing bacteria (Clostridium botulinum)	<10	1.2 10 <sup>2</sup>	0.7 10 <sup>2</sup>
Salmonella spp.	-	-	-

497 Results are expressed as ufc/g; the symbol "-" indicates that the organism was not present

498 **Table 4.** Physical and chemical parameters on Cinta Sense dry-fermented sausages manufactured

499 with natural antioxidant as replacement of sodium nitrite.

	GSE	CHE	NIT	<b>SEM</b> <sup>a</sup>	Pb
pH 0 days	6.13	6.40	6.35	0.05	n.s.
pH 24 days	6.02b	6.04b	6.10a	0.04	**
aw	0.89	0.88	0.90	0.01	n.s.
L*	43.21	41.67	42.37	0.67	n.s.
a*	17.22a	15.92b	18.06a	0.41	**
b*	5.31b	4.87b	6.48a	0.21	**
Weight loss (%)	40.23	43.30	42.92	1.35	n.s.
Moisture (%)	30.43	29.97	30.94	0.66	n.s.
Fat (g/100 g dm)	41.82	41.01	41.32	0.33	n.s.
Protein (g/100 g dm)	48.88	49.74	49.14	0.31	n.s.
Ash (g/100g dm)	8.46	8.69	8.80	0.12	n.s.
TBARs (mg MDA/kg)	1.05	0.98	0.93	0.05	n.s.
SFA (mg/100g)	8.05	7.57	7.71	0.23	n.s.
MUFA (mg/100g)	9.72	9.17	9.10	0.25	n.s.
PUFA (mg/100g)	3.32	3.38	3.47	0.10	n.s.

500 Values are reported as means of the two replications within the same treatment, where GSE is grape 501 seed extract added group, CHE the chestnut extract added group and NIT the control group added 502 with sodium nitrite

<sup>a</sup> Standard error

<sup>b</sup>P value of natural antioxidant effect \*\* p<0.01, \* p<0.05, different letters in the same row indicate</li>
significant differences at p<0.05.</li>

506

**Table 5.** Texture traits of Cinta Sense dry-fermented sausages manufactured with natural
antioxidant as replacement of sodium nitrite.

	GSE	CHE	NIT	<b>SEM</b> <sup>a</sup>	Pb
Hardness (N)	104.93	102.82	102.68	6.38	n.s.
Cohesiveness	0.38ab	0.42a	0.35b	0.01	**
Gumminess	39.57	42.93	35.99	2.11	n.s.
Springiness	3.04ab	3.26a	2.70b	0.13	*
Chewiness (N)	120.67ab	139.55a	96.79b	8.36	**

509 Values are reported as means of the two replications within the same treatment, where GSE is grape 510 seed extract added group, CHE the chestnut extract added group and NIT the control group added 511 with sodium nitrite

- 512 <sup>a</sup>Standard error
- <sup>b</sup>P value of natural antioxidant effect \*\* p<0.01, \* p<0.05, different letters in the same row indicate

514 significant differences at p < 0.05.

<sup>Table 6. Volatile compounds in Cinta Sense dry-fermented sausages manufactured with natural
antioxidant as replacement of sodium nitrite.</sup> 

Compound	LRI <sup>a</sup>	RI <sup>b</sup>	GSE	CHE	NIT	SEM <sup>c</sup>	
Lipid auto-oxidation							
Pentane	500	а	0.59	0.60	0.56	0.19	
Propanal	524	а	0.19	0.13	0.13	0.06	
Isopropanol (45)	539	-	1.20	0.79	1.13	0.40	
Hexane	600	а	0.76	0.74	0.56	0.22	
1-Propanol	613	а	4.90	4.90	3.72	2.37	
Octane	800	а	4.20	3.498	3.90	0.98	
Propanoic acid (74)	814	а	0.14a	0.05b	0.02b	0.05	
1-Pentanol	827	а	0.34ab	0.43a	0.22b	0.16	
Hexanal	841	a	2.62a	2.27 a	1.36 b	0.27	
1-Hexanol	924	а	0.79	0.75	0.88	0.16	
Decane	1000	а	1.10a	1.26a	0.48b	0.18	
Dodecane	1200	а	0.96	1.14	0.92	0.37	
Tridecane	1300	а	0.36	0.44	0.34	0.10	
Tetradecane	1400	а	0.18b	0.87a	0.22b	0.16	
Total			19.79a	16.19ab	14.57b	0.96	
Spices							
α-Thujene	934	b	24.40	26.07	27.29	6.22	
α -Pinene	940	a	17.84	19.00	17.33	3.68	
Sabinene+ <sub>β</sub> -pinene	986	a	133.28	149.70	145.97	19.83	
β-Myrcene	1003	a	37.19b	53.47a	39.01b	9.25	
α-Phellandrene (93)	1019	b	8.71b	12.39a	9.41b	2.41	

3-Carene	1023	а	73.20	86.24	77.49	15.77	
α -Terpinene	1035	а	3.66b	4.54a	3.28b	0.78	*
Unknown (57)	1042	b	0.17	0.16	0.14	0.11	
Limonene	1046	а	307.31b	401.19a	289.74b	60.75	**
β-Phellandrene (93)	1051	b	13.98b	19.28a	14.42b	3.74	*
p-Cymene (119)	1052	b	8.49	9.078	8.98	2.73	
β-Ocimene	1067	b	1.18b	1.87a	1.15b	0.44	**
4-Carene	1073	b	0.21	0.27	0.22	0.12	
Unkwon	1075	b	3.21b	4.28a	3.16b	0.93	*
Styrene	1091	a	2.20	2.43	2.34	0.31	
Terpene	1099	b	0.95	1.46	1.35	0.31	
Terpinolene	1101	b	4.64b	6.48a	4.46b	1.20	**
Unknown (93)	1120	b	0.51	0.55	0.56	0.11	
Linalool (93)	1150	a	0.28b	0.33a	0.32ab	0.04	*
$\beta$ -Terpinene/ $\gamma$ -Terpinene	1158	b	1.4	1.37	1.43	0.24	
4-Terpineol	1231	а	2.14	2.31	2.12	0.29	
Estragole	1249	а	1.28	1.33	1.20	0.21	
α -terpineol	1256	а	0.82	0.91	0.88	0.13	
δ-Elemene	1342	b	3.58a	4.19a	2.74b	0.66	**
α-Cubebene	1348	b	0.85a	0.92a	0.68b	0.12	**
Cyclosativene (161)	1402	b	0.02b	0.04a	0.02b	0.00	*
Copaene (161)	1406	b	1.50a	1.42a	1.16b	0.28	*
β-Cubabene (161)	1421	b	0.20a	0.19a	0.16b	0.04	*
β-Elemene (93)	1423	b	0.11b	0.13a	0.09b	0.04	**
$\alpha$ -Bergamotene	1433	b	0.79a	0.91a	0.69c	0.13	**

trans-α-Bergamotene	1450	b	0.67	0.84	0.76	0.18	
β-Caryophyllene	1455	b	107.97a	113.01a	91.99b	13.04	**
$\alpha$ -Caryophyllene	1486	b	4.87a	4.99a	4.19b	0.68	*
Isocaryophillene	1498	b	0.65	0.84	0.57	0.26	
Isolongifolene	1510	b	10.64b	11.05b	25.91a	10.05	**
Valencene	1513	b	0.00b	5.31a	0.00b	1.78	**
γ-Cadinene	1518	b	1.72b	1.89b	2.92a	0.56	**
δ-Cadinene	1529	b	0.96b	1.04b	1.29a	0.20	**
Total			783.55b	984.05a	785.83b	134.67	**
Bacterial metabolism							
Lipid β-oxidation							
2-Pentanone	734	а	5.64	5.96	6.92	1.79	
3-Octanone	1032	a	2.23	2.51	2.63	0.65	
Total			7.87b	8.47ab	9.55a	0.63	*
Esterase activity							
3-Methyl-1-butanol acetate	907	a	0.59	0.49	0.51	0.12	
Carbohydrate							
fermentation							
Acetaldehyde	462	а	1.26a	0.86b	0.89b	0.25	**
Ethyl alcohol	507	а	6.79a	5.65ab	3.68b	2.23	*
Acetone	529	а	72.92	76.47	72.72	20.58	

2,3-Butanedione	627	а	9.76a	7.47b	7.83b	2.01	*
2-Butanone	632	а	241.96	287.47	332.22	91.70	
2-Butanol	644	a	19.99	20.52	29.19	11.02	
Acetic acid (60)	720	a	11.99	12.19	7.08	5.34	
1-Butanol (56)	727	a	0.28	0.18	0.71	0.67	
2-Pentanol	756	a	0.49	0.43	0.38	0.15	
3-Hydroxy-2-butanone	782	a	124.14a	89.35b	136.91a	31.84	**
2,3-Butanediol (45)	884	a	6.20a	1.50b	3.76b	2.92	**
2,3-Butanediol (45)	892	a	4.17	2.30	2.36	1.98	
Butanoic acid (60)	894	а	0.52a	0.49a	0.18b	0.254808	*
Total			499.97b	489.85b	597.61a	23.69	**
Amino acid degradation							
2-Methylpropanal	595	а	0.54b	0.66a	0.39b	0.17	**
3-Methylbutanal	691	а	11.34	12.43	15.34	4.53	
2-Methylbutanal	701	a	6.18	7.53	8.85	2.88	
Toluene	789	а	51.99	56.94	59.18	6.88	
3-Methylbutanol	795	a	7.39b	8.46b	15.13a	5.92	*
2-Methylbutanol	797	a	1.27	1.44	1.82	5.92	
Pyrrole	845	a	0.31	0.30	0.25	0.05	
2-Methylpropanoic acid	868	a	2.01	1.70	2.70	1.06	
Ethylbenzene (91)	884	а	0.09b	0.18a	0.09b	0.05	**
3-Methylbutanoic acid (60)	942	а	2.52	2.25	2.27	1.05	
2,5- Dimethylpirazine (108)	943	a	0.34	0.34	0.36	0.09	
2-Methylbutanoic acid	948	a	2.32	2.73	2.98	1.25	

2-Acetyl-1-pyrroline	961	a	0.72b	0.91b	1.37a	0.37	**
Methional	986	a	0.16	0.18	0.15	0.02	
Benzaldehyde (106)	1020	a	1.14	1.40	1.27	0.33	
Benzeneacetaldehyde	1110	a	6.61a	4.14b	2.72b	3.17	*
Tetramethylpyrazine	1118	a	0.03	0.02	0.03	0.01	
Benzylalcohol (79)	1122	a	0.27a	0.18b	0.19b	0.06	**
Phenylethyl alcohol (91)	1195	a	0.53	0.64	0.81	0.35	
Total			96.59b	103.09ab	115.63a	4.55	*
Total microbial metabolism			605.04b	601.91b	723.03a	24.32	**
Unknown or contaminant compounds							
Carbon disulfide (76)	537	a	5.80	6.42	6.70	1.89	
p-Xylene (91)	892	а	0.09b	0.31a	0.11a	0.09	**
2-Butoxyethanol	953	a	2.45	2.23	2.33	0.33	
4-Methylphenol (108)	1199	а	0.10	0.06	0.08	0.03	

517 Values are reported as means of the two replications within the same treatment, where GSE is grape 518 seed extract added group, CHE the chestnut extract added group and NIT the control group added 519 with sodium nitrite.

520 Abundance expressed as AU x  $10^{-6}$  (AU: abundance unit, expressed as total ion chromatogram 521 (TIC) or area of the target ion shown in parenthesis).

<sup>a</sup>Linear retention indices (LRI) of the compounds eluted from the GC-MS using a DB-624 capillary
 column.

- <sup>b</sup>Reliability of identification: a, identification by mass-spectrum and by coincidence with the LRI of
- an authentic standard; b, tentatively identification by mass-spectrum.
- 526 <sup>c</sup>Standard error
- <sup>d</sup>P value of natural antioxidant effect \*\* p<0.01, \* p<0.05, different letters in the same row indicate
- 528 significant differences at p < 0.05.

- **Table 7.** Odor active compounds identified by GC-O in Cinta Sense dry-fermented sausages manufactured with natural antioxidant as replacement
- 530 of sodium nitrite.

Compound	GC-MS <sup>1</sup>		GC-O <sup>2</sup>				
	LRI	LRI std	LRI initial	LRI final	RI <sup>3</sup>	Odour Description	<b>DF</b> 4
Lipid auto-oxidation							
1-Propanol	613	611	613	618	а	vegetal, green, pungent, fresh, floral,	5
Propanoic Acid	814	806	800	810	а	tasty, fresh, green, cheese, cured, pungent	4
1-Pentanol	827	823	821	827	а	vegetable, pungent, unpleasant, cabbage, acid	6
Hexanal	841	839	834	844	а	green, grass, vegetable, fresh	8
Spices							
Linalool	1150	1145	1141	1149	а	fresh, floral, cabbage, unpleasant, soap	7
$\beta$ -Terpinene/ $\gamma$ -Terpinene	1158		1160	1167	b	cooked, cooked vegetable, floral, pungent, resin	8
β-Myrcene	1003	1003	1000	1004	a	irritating, spicy, pepper, green, leaves, earthy	10
α-Terpinene	1035	1035	1030	1034	а	mushrooms, wetness, burnt, unpleasant, pungent, pine, woody, earthy	12
Unkown terpene	1075		1076	1080	c	earthy, green, vegetable, fresh, fruity, cologne	5
Terpinolene	1101	1106	1107	1113	а	floral, rose, grass, green	11

α-Thujene	934		929	933	c	sour, vinager, unpleasant, fruity,	5
Lipid β-oxidation							
2-Pentanone	735	731	725	733	a	floral, green, fresh, oxidized fat, cheese	8
Esterase activity							
3-Methyl-1-butanol acetate	907	905	905	911	a	sweet, fresh, floral	7
Carbohydrate fermntation							
2-Butanone	632	629	631	642	a	sweet, slightly unpleasant, green, grass	5
2-Butanol	644	643	650	661	a	sweet, caramel, malt, unpleasant,	5
Acetic acid	720	718	714	722	a	grass, vegetable, fresh, wine, green	5
Amino acid degradation							
2-Methylpropanal	595	590	602	609	a	acid, floral, green, weak	4
3-Methylbutanal	691	687	690	697	a	caramel, chocolate, grass, fresh	6
2-Methylbutanal	702	699	691	700	а	sweet, floral, fruity, toasted,	6
3-Methylbutanol	795	793	791	793	а	sweet, spicy, toasted, floral	4
2-Methylpropanoic acid	868	864	869	872	a	cheese, roasted, cured, green, slightly sweet, unpleasant	6

Ethylbenzene	884	881	884	891		a	earthy, fresh, green, mushroom	4
3-Methylbutanoic acid	942	941	922	926		a	cheese, rancid, oxidized fat	8
2,5-Dimethylpyrazine	943	943	936	943		a	meaty, cooked potatoes, sweet, buttery	7
2-Acetyl-1-pyrroline	961	960	960	964		a	roasted, nuts, bread, pop-corn, biscuits, fried potatoes	12
Methional	986	964	966	969		a	mashed potato, cooked onion, roasted meat	9
Tetramethylpyrazine	1118	1118	1115	1121		a	earthy, green, grass, wetness, fresh	7
Unknown or contaminant compounds								
Carbon Disulfide	537	537	531	543		a	weak, burnt, malt	4
Unknown		776	762	766		c	cured, meat, acid, fresh, acid	6
Unknown	1190	1182	1176	1182		c	roasted, fried nuts, biscuits	11

<sup>1</sup>Linear retention index (LRI) of the compounds eluted from the GC-MS and LRI of standard compound

<sup>532</sup> <sup>2</sup>Initial and end linear retention index of aroma compound in GC-FID-O

<sup>3</sup>Reliability of identification (RI): a, identification by mass spectrum, coincidence with LRI of an authentic standard and by coincidence of the

- assessor's descriptors with those described in the Fenaroli's handbook of flavor ingredients (Burdock, 2002); b, tentatively identification by mass
- 535 spectrum; c, unknown compounds.
- <sup>4</sup>Detection frequency value.