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# 1 Can sensory boar taint levels be explained by fatty acid 2 composition and emitted volatile organic compounds in addition to 3 androstenone and skatole content?

4 Clément Burgeon<sup>a,\*</sup>, Maria Font-i-Furnols<sup>b</sup>, María Dolores Garrido<sup>c</sup>, María Belén Linares<sup>c</sup>, Yves  
5 Brostaux<sup>d</sup>, Gerard Sabeña<sup>b</sup>, Marie-Laure Fauconnier<sup>a,†</sup>, Núria Panella-Riera<sup>b,†</sup>

6 <sup>a</sup> Laboratory of Chemistry of Natural Molecules, Gembloux Agro-Bio Tech, Université de Liège,  
7 Passage des Déportés 2, 5030 Gembloux, Belgium

8 <sup>b</sup> IRTA-Product Quality, Finca Camps i Armet, 17121 Monells, Spain

9 <sup>c</sup> Department of Food Science and Technology, Veterinary Faculty, University of Murcia, 30071  
10 Espinardo Murcia, Spain

11 <sup>d</sup> Statistics, Informatics, and Applied Modeling Unit, Department of AgroBioChem, Gembloux Agro-  
12 Bio Tech, Université de Liège, Passage des Déportés 2, 5030 Gembloux, Belgium

## 13 Abstract:

14 This study aimed at understanding which molecules were responsible for the differences existing in  
15 boar taint sensory evaluation. The latter was therefore linked to the results of skatole and androstenone  
16 chemical analyses, fatty acid composition and VOC profiles of heated backfat. This study confirmed  
17 that some discrepancy exists between chemical analysis and sensory evaluation of tainted backfats.  
18 Significant correlations between human nose scores and fatty acid composition were not revealed.  
19 Strong correlations between emissions and contents in skatole and androstenone were found. Oxidation  
20 products of polyunsaturated fatty acids, with fatty odor descriptors, were found to be more present in  
21 the VOC profiles of boar fat considered untainted through the human nose methodology. Weak  
22 coefficient of determination for partial least square regression indicates that other factors, yet unknown,  
23 are responsible for sensory evaluation outcomes. These findings hence support the idea that high human  
24 nose score is mainly due to boar taint compounds rather than general differences in VOC profiles.

25 **Keywords:** boar taint, androstenone, skatole, human nose, fatty acid, VOC

26 \* Corresponding author: *E-mail address:* [cburgeon@uliege.be](mailto:cburgeon@uliege.be) (Clément Burgeon)

27 † These authors have contributed equally to this work and share last authorship

## 28 **1. Introduction**

29 Surgical castration of piglets without anesthesia and analgesia is the most common practice to ensure  
30 the absence of boar taint (Fredriksen et al., 2009), a strong smell created by a variety of molecules  
31 released upon cooking of pork meat (Bonneau et al., 2000) and perceived as unpleasant by 14.3-41.0%  
32 of European pork consumers (Blanch et al., 2012). However, due to growing animal welfare concerns  
33 in the European Union, there have been some actions, such as the Noordwijk Declaration (2007),  
34 the Düsseldorf Declaration (2008) and the Brussels Declaration (2010), to reduce or eliminate this  
35 practice. As a result, Norway banned surgical castration without pain relief in 2002, Switzerland banned  
36 it in 2016, Sweden in 2016, Germany at the end of 2020 and lastly, France banned surgical castration  
37 without anesthesia at the end of 2021 (De Briyne et al., 2016; Lin-Schilstra & Ingenbleek, 2021).  
38 Currently, three viable alternatives exist: surgical castration with pain reliefs, immunocastration and  
39 finally raising entire males (Bonneau & Weiler, 2019). A main disadvantage of this last option remains  
40 that high levels of boar taint are found in 5-10% of the cases (Borrisser-Pairó et al., 2016) and its  
41 occurrence could be even higher depending on the pig's characteristics and environmental factors  
42 (Aluwé et al., 2020). On the other hand, although immunocastration has been found to be overall very  
43 efficient, some non-responders might still occur, with the potential accumulation of boar taint (Čandek-  
44 Potokar et al., 2017; Font-i-Furnols et al., 2012; Kress et al., 2020).

45 Being able to correctly discriminate tainted from untainted pigs is therefore a topic of current interest.  
46 Throughout the years, many methods have been developed with the aim of offering rapid, cheap and  
47 reliable boar taint detection for slaughterhouses and cutting plants. Amongst these techniques, laser  
48 diode thermal desorption-tandem mass spectrometry (LDTD-MS/MS) is a recently developed method  
49 that focuses on the specific detection and quantification of skatole and androstenone (Lund et al., 2021)  
50 and is currently being tested in Danish slaughterhouse (Burgeon, Debliquy, et al., 2021; Font-i-Furnols  
51 et al., 2020). These two molecules are well known to be the two major contributors to boar taint which  
52 give respectively a strong fecal and urine smell to pork meat (Patterson, 1968; Vold, 1970). Yet, only  
53 50% of the variation in boar taint is due to the combination of skatole and androstenone (Hansson et  
54 al., 1980) and the importance of other compounds to boar taint has been pointed out in several studies  
55 on sensory evaluation (Font-i-Furnols et al., 2008; Mathur et al., 2012; Trautmann et al., 2016;  
56 Whittington et al., 2011).

57 A variety of molecules have therefore been suggested to potentially contribute to boar taint. The origin  
58 of each molecule and their contribution to the overall boar taint is variable. In fact, some are derived  
59 from skatole such as indole and 2-aminoacetophenone, some are related to androstenone, such as 3 $\alpha$ -  
60 androstenol and 3 $\beta$ -androstenol and others have synthesis pathways which are less clear or nonetheless  
61 not directly linked to skatole and androstenone. The latter include *p*-cresol, 4-ethylphenol found in boar

62 preputial fluid and 1,4-dichlorobenzene found in boar fat, but which occurrence in boar fat could be due  
63 to external contamination (Brooks & Pearson, 1989; Fischer et al., 2014; Garcia-Regueiro & Diaz,  
64 1989; Patterson, 1967; Solé & Regueiro, 2001; Watson & Patterson, 1982).

65 Many of the above-mentioned molecules are found in trace amounts in boar fat samples. Therefore,  
66 although they might have an unpleasant odor descriptor and a low odor threshold, which allows them  
67 to contribute to the overall boar taint, other compounds are found in much greater concentration in the  
68 headspace of heated fat, as pointed it out in a previous study by Burgeon, Markey, et al. (2021).  
69 Consequently, even without being directly related to the urine and fecal smell of boar taint, the latter  
70 still contribute to the overall smell perceived during sensory analysis. Many of these molecules, such  
71 as free fatty acids and aldehydes, originated from the oxidation of lipids starting at 70°C. The presence  
72 of such molecules in the headspace of heated fat is explained by the fact that skatole and androstenone  
73 are lipophilic molecules with low vapor pressure ( $7.3 \times 10^{-4}$  kPa and  $1.3 \times 10^{-6}$  kPa at 25 °C,  
74 respectively) and need to be heated at high temperatures to be released and detected by the human nose  
75 (Burgeon, Markey, et al., 2021).

76 Human nose detection (*i.e.* sensory analysis) remains a method of choice given its ease of  
77 implementation, low cost per analysis and satisfactory performances results. Additionally, it is to date  
78 the only method currently in use, which takes into account all of the volatile organic compounds (VOCs)  
79 constituting the complex smell of boar taint. This implies that this method could be the only one able  
80 to detect samples which are considered tainted, although chemical analysis would classify them as  
81 untainted given skatole and androstenone concentrations below rejection thresholds.

82 Research has already been performed on fatty acid composition of backfat from pigs of different sex,  
83 weight and breed (Font-i-Furnols et al., 2019; Raj et al., 2010). Research on fatty acid composition of  
84 backfat with various boar taint levels, and its relationship with skatole and androstenone content as well  
85 as sensory analysis has also been reported (Liu et al., 2017; Mörlein & Tholen, 2015). Lastly, the  
86 relationship between fatty acid composition and VOC emission of lard was studied, without bringing  
87 attention to skatole and androstenone content of the analyzed meat cut, as this was not the aim (Serra et  
88 al., 2014).

89 In this study, the authors consider all four aspects mentioned above: (i) fatty acid composition of boar  
90 fats with varying boar taint levels, (ii) sensory evaluation of boar taint, (iii) chemical levels of boar taint  
91 compounds, androstenone and skatole, and (iv) the VOCs emitted when heating backfat. Through the  
92 comprehension of the correlation between these variables and the human nose score attributed during  
93 sensory evaluation, this work aims to understand which factor, other than skatole and androstenone  
94 content could explain the human nose scores. In other words, this should clarify the source of existing  
95 discrepancies between backfat classification based on skatole and androstenone content analysis and  
96 sensory evaluation.

## 97 **2. Material and methods**

### 98 *2.1. Samples*

99 Backfat samples from 30 boars were randomly selected from a sample of 106 boars at the  
100 slaughterhouse. These were sampled during different days and from animals reared under different  
101 production systems with different managing and feeding strategies. To ensure the presence of boar  
102 tainted carcasses, one trained panelist smelt the subcutaneous fat of the carcass online, close to the neck,  
103 after heating the fat with a gas-powered torch heated plate (human nose method). Carcasses classified  
104 as boar tainted were selected together with some untainted samples, to ensure enough variability.  
105 Approximately 30g of fat from the neck of the selected carcasses was collected and immediately frozen  
106 at -20°C until further analysis (maximum storage time of 1 year). From all the samples, fatty acid  
107 composition and skatole and androstenone were chemically analyzed as described in section 2.3 and  
108 2.4, respectively. The backfat samples were selected in order to fit into four categories based on their  
109 skatole and androstenone content and used in further analysis. Samples with low and high skatole  
110 content were distinguished using the threshold value of 0.2 µg/g liquid fat. This value has been used in  
111 other works (Bonneau et al., 2000; Burgeon et al., 2021) although skatole was measured in fat tissue,  
112 which provides lower values than those obtained in liquid fat (Haugen et al., 2012). Nevertheless, it  
113 was difficult to find enough samples with high levels of skatole (and low androstenone) within the pre-  
114 selected fat, confirming what was suggested by Zamaratskaia & Squires (2009). Low and medium  
115 androstenone content were separated by a threshold value of approximately 1.0 µg/g liquid fat, while  
116 medium and high androstenone concentrations were hence distinguished by a value of approximately  
117 1.5 µg/g liquid fat. These thresholds have been chosen to ensure variability of androstenone levels,  
118 since the global samples were highly variable for this compound. According to this, the four groups  
119 created were: low skatole, low androstenone (LS/LA, n=8); low skatole, high androstenone (LS/HA,  
120 n=7); high skatole, medium androstenone (HS/MA, n=7); high skatole, high androstenone (HS/HA,  
121 n=8).

### 122 *2.2. Sensory evaluation*

123 Sensory evaluation of boar taint, also known as human nose, was performed, at line, by three  
124 trained panelists on the 30 selected carcasses. Each sample was evaluated in duplicate. Subcutaneous  
125 fat samples were heated with a soldering iron (Soldering iron station Analogue 58W, 150-450°C,  
126 Basetech, Austria) at approximately 250°C until the surface of the fat melted. Immediately, fat samples  
127 were smelt by the three trained panelists (three androstenone and skatole sensitive women between 40  
128 and 55 years old) and classified according to a 4 points scale: 0: no boar taint; 1: weak boar taint; 2:  
129 moderate boar taint; 3: strong boar taint. Each sample was evaluated twice. The average human nose  
130 score (HNS) was obtained and used for further analysis. The global panel performance, evaluated with  
131 fat samples with known levels of androstenone and skatole, was sensitivity= 0.74; specificity= 0.70;

132 accuracy= 0.74. Individual panelist performance was characterized and the Positive Predictive Value  
133 (PPV) as well as the Negative Predictive Value (NPV) were calculated (Appendix A). Detection  
134 thresholds evaluated using smell strips (Meier-Dinkel, Trautmann, et al., 2013) were 0.2 µg/g for  
135 androstenone and 0.05 µg/g for skatole.

### 136 *2.3. Fatty acid composition*

137 For the FA analysis, a portion of 10 g of frozen fat was thawed at 4°C during 24 hours. Fat was ground  
138 with a commercial grinder, and FA were quantified as FAME (FA methyl ester) using 25 mL of sodium  
139 methylate and 30mL paratoluensulfonic acid for transesterification. The sample was methylated by  
140 incubation in a sand bath at 80 °C for 50 min. For FAME solubilization 20 mL of heptane was used. An  
141 aliquot of 0.4 µL was introduced by split injection into a capillary column (60 m x ID 0.25 mm, Agilent  
142 HP88; 0.25-µm film thickness, Barcelona, Spain). Helium was the carrier gas at 1.5 mL/min. Column  
143 temperature was initially at 140°C for 5 min, was increased by 4°C/min to 240°C and maintained for 20  
144 min at this temperature. Individual FAME were identified by retention time with reference to Supelco®  
145 37 Component FAME Mix (47885-U Sigma Chemical Co., St. Louis, MO). Data is expressed in mg/100  
146 g of adipose tissue.

### 147 *2.4. Skatole and androstenone quantification in backfat*

148 Quantification of skatole and androstenone in backfat were performed by stable isotope dilution  
149 analysis – headspace solid-phase microextraction – gas chromatography/mass spectrometry (SIDA-HS-  
150 SPME-GC/MS) developed by Fischer et al. (2011). Results are expressed as µg/g of liquid fat.

151 Briefly, thawed and skinless backfat samples are diced and microwave heated for 2 min at 180W. After  
152 separation of connective tissue and liquid fat, 500 mg of fat was transferred into a 2 mL plastic vial and  
153 spiked with 250 ng of androstenone-d<sub>3</sub> and 50 ng of skatole-d<sub>3</sub> to achieve final concentrations of 500  
154 ng/g androstenone-d<sub>3</sub> and 100 ng/g skatole-d<sub>3</sub>. The sealed plastic vial was then shaken for 30s, stored  
155 for 10 min at 55°C and mixed again for 30 s in order to allow for equilibration of the standards.  
156 Extraction of the compounds of interest was then performed by adding 1 mL of methanol and then  
157 shaking 30s, heating for 10 min at 55°C and shaking again 30s. The samples were then centrifuged (10  
158 min, 6500 rpm, -15°C). The methanolic supernatant was transferred into a 10 mL headspace vial and  
159 evaporated at 40°C by a gentle stream of air. Once dryness was achieved, the vial was sealed and placed  
160 in an autosampler device (Varian Combi Pal, Darmstadt, Germany), operating with a heated agitator  
161 and an SPME assembly. HS-SPME was carried out as follows: equilibration for 5 min at 100°C;  
162 extraction for 30 min at 100°C with a fused-silica fiber coated with 65 µm  
163 poly(dimethylsiloxane)/divinylbenzene (PDMS/DVB) (Supelco, Bellefonte, PA); desorption for 20  
164 min within the injector.

165 A GC-MS (GC-450 – MS-240 ion trap, Varian, Darmstadt, Germany) equipped with a Varian VF-5ms  
166 capillary column (30 m × 250 μm × 0.25 μm, Varian, Darmstadt, Germany) was used for the analyses.  
167 Helium was used as a carrier gas at a flow rate of 1.0 mL/min. The oven temperature program was set  
168 as follows: start at 50°C, hold for 3 min, then raise to 160°C at a rate of 10°C/min, followed by a raise  
169 of 5°C/min up to 240°C, hold for 1 min. Injection was performed at 270°C in splitless mode for 3 min,  
170 the split valve was then opened in a split ratio 1:100.

171 Mass spectrometry data was obtained with a full scan acquisition mode using electron impact ionization  
172 (EI). The peak area ratios of analyte and internal standard (IS) were later determined by displaying the  
173 specific mass fragments of each analyte and its corresponding IS in selected ion monitoring (SIM)  
174 mode. The selected mass traces (m/z) were as follows: skatole m/z 130, skatole-d<sub>3</sub> m/z 133 + 134 and  
175 androstenone m/z 257 + 272 and androstenone-d<sub>3</sub> m/z 260 + 275.

## 176 *2.5. Analysis of VOCs generated through heating of backfat*

177 The analysis of VOC profiles produced through the heating of boar backfat was performed following  
178 incubation of fat at 150°C and analysis by SPME-GC-MS according to the method described by  
179 Burgeon, Markey, et al. (2021).

180 Briefly, 2.5g of backfat was cut and cooled with liquid nitrogen. The sample was then ground for 5s  
181 with an analytical grinder (A11 basic analytical mill, IKA) and a 1.0g of sample was recovered in a vial  
182 and stored at -20°C until analysis. The next step of this method consisted in a 20 min incubation at  
183 150°C in a heated agitator (Gerstel, Mülheim an der Ruhr, Germany). VOCs sampling was then  
184 achieved by exposing a divinylbenzene/carboxen/polydimethylsiloxane (DVB/CAR/PDMS, 50/30 μm)  
185 SPME fiber (Supelco, Darmstadt, Germany) to the headspace for 5 min. The vials were shaken at 250  
186 rpm (agitator on/off time: 10 s/1 s) during incubation and extraction. Fiber desorption took place for 2  
187 min. Injection was performed in splitless mode at 270°C. The fiber was left for 20 min at injection  
188 temperature for reconditioning.

189 A GC-MS (7890A-5975C, Agilent Technologies, Santa Clara, CA, USA) equipped with an HP-5 MS  
190 capillary column (30 m × 250 μm × 0.25 μm, Agilent Technologies, Santa Clara, CA, USA) was used  
191 for the analyses. Helium was used as a carrier gas at a flow rate of 1.2 mL/min. The oven temperature  
192 program was set as follows: start at 40°C, hold for 3 min, then raise to 300°C at a rate of 5°C/min, hold  
193 for 5 min at 300°C. The mass spectrometer was set to have a temperature of 230°C at the ion source  
194 and 150°C at the quadrupole. Mass spectrometry data was obtained with a SIM/SCAN acquisition  
195 mode. In SIM mode, the targeted ions were: m/z 130 for skatole and m/z 257, and 272 for androstenone.  
196 The peak area of these ions (expressed in atomic mass unit, amu) was integrated to study the relationship  
197 between content and emissions as described later, ie. these peak areas constitute the “emission data”.

198 In SCAN mode, mass spectra were scanned from 35 to 500 amu. Component identification was then  
199 performed by comparison of the obtained spectra with reference spectra from the NIST17 database  
200 (National Institute of Standards and Technology, Gaithersburg, USA). Additionally, experimental  
201 retention indices (RI) were calculated following the injection of a n-alkanes C8-C30 mixture (Sigma  
202 Aldrich, Darmstadt, Germany) under the same chromatographic conditions as those previously  
203 mentioned. This allowed the comparison of these RI with literature RIs obtained from the NIST Mass  
204 Spectrometry Data center.

205 Lastly, pure standards were injected for skatole (CAS n° 83-34-1, Sigma Aldrich) and androstenone  
206 (CAS n° 18339-16-7, Sigma Aldrich, Darmstadt, Germany) to ensure that the peaks were correctly  
207 identified.

208 The only difference to the method described in Burgeon, Markey, et al. (2021) is that the incubation  
209 temperature was fixed to 150°C and no internal standard was added for semi-quantification (of skatole  
210 and androstenone) as this was not the aim of the article.

## 211 *2.6. Data analysis*

212 Pearson correlation coefficients were determined between fatty acids and HNS, between HNS and  
213 androstenone and skatole content and between the skatole and androstenone emission and content data.  
214 Fatty acid compositions of the different chemical classification groups (Appendix B), were analyzed  
215 using one-way ANOVA (one fixed factor: taint group). Volatile organic compounds and fatty acid  
216 compositions of the different human nose (HN) classification categories were also analyzed using one-  
217 way ANOVA (one fixed factor: HN category). When the means were significantly different ( $p < 0.05$ ),  
218 a Tukey-Kramer comparison test was performed. Both Pearson correlation coefficients and ANOVA  
219 were established with Minitab 19 software (Minitab Inc., State College, PA, USA).

220 Principal components analyses were performed on fatty acid composition data and VOC data to detect  
221 existing trends between the different samples. A partial least square regression (PLSR) was used to  
222 develop a mathematical model trying to predict human nose score (HNS) by taking into account skatole  
223 and androstenone content, fatty acid content and VOC emission. A cross-validation was performed to  
224 choose the optimal number of PLS components. The number of components chosen for the model were  
225 those that yielded in the lowest cross-validated root mean square of prediction (RSMEP) and highest  
226 coefficient of determination ( $R^2$ ). The variable importance in projection (VIP) scores were then  
227 analyzed. Both PCA and PLSR were conducted in R (R 4.0.2 software, R Development Core Team,  
228 Boston United States).

229 PCA individual plots were performed in R. All other graphs and tables were established on Excel  
230 (Microsoft Office 2016).



### 231 3. Results and discussion

#### 232 3.1. Agreement between human nose evaluation and chemical analysis

233 As a reminder, based on the results obtained through chemical analysis, four groups were created: low  
234 skatole, low androstenone (LS/LA, n=8); low skatole, high androstenone (LS/HA, n=7); high skatole,  
235 medium androstenone (HS/MA, n=7); high skatole, high androstenone (HS/HA, n=8). The samples'  
236 taint was also assessed through sensory evaluation. The results obtained for both tests (chemical  
237 analysis and human nose evaluation) have been compared and are presented in **¡Error! No se encuentra  
238 el origen de la referencia..**

239 It can be noticed from Figure 1 that almost all fat samples scored as untainted through the human nose  
240 method are found in the LS/LA part of the graph (6 out of the 7 with HNS=0). Similarly, most of the  
241 fats graded as highly tainted (3 out of the 4 fats with HNS=3) are found in the HS/HA region. In this  
242 class there are also 4 samples with  $HNS \geq 2$  and one fat with HNS=1. There seems to be positive  
243 correlation in this region between the two methods – when skatole and androstenone content increases,  
244 the HNS increases accordingly. These observations support the idea that the human nose is an overall  
245 good representation of the chemical analyses available.

246 The rest of the fat samples with intermediate HNS are found in analytical groups combining a high level  
247 of skatole or androstenone and a lower level of androstenone or skatole. Interestingly, only fats with  
248 HNS greater than 1.5 are found in the HS/MA region when most of the samples in the LS/HA region  
249 have HNS lower or equal to 1.5 (5 out of 7 fats found in this region). This result shows a greater  
250 influence of skatole level in the HNS which can also be seen when the average HNS by class group is  
251 studied (Appendix C). This is in agreement with the results presented by Mörlein et al. (2016).

252 The higher difficulty to evaluate androstenone than skatole in the boar taint perception carried out with  
253 trained panelist is in agreement with previous works (Aluwé et al., 2022; Dijksterhuis et al., 2000; Font  
254 et al., 2009; Lunde et al., 2010). In this regard, HNS is higher and similar in HS/HA and HS/MA group  
255 ( $2.38 \pm 0.69$  and  $2.36 \pm 0.48$ , respectively, both with high levels of skatole), is intermediate in the LS/HA  
256 class ( $1.29 \pm 0.95$ , with low levels of skatole) and is low in the LS/LA class ( $0.31 \pm 0.70$ ; with low levels  
257 of skatole). Having high levels of androstenone classified differently with the HNS points out the  
258 difficulty of this sensory classification, especially when skatole is absent. This can increase the false  
259 negative if androstenone levels are high and skatole are low, which agrees with the higher false negative  
260 with high androstenone and low skatole levels reported by Meier-Dinkel et al. (2015). This agrees with  
261 the higher correlation between HNS and skatole (0.68,  $P < 0.001$ ) compared to those between HNS and  
262 androstenone (0.37,  $P = 0.042$ ), that are in line with those reported by Mathur et al. (2012), 0.69 between  
263 HNS and skatole and 0.42 between HNS and androstenone. In general, these intermediate levels are  
264 more difficult to be detected with the sensory evaluation.

265 Several hypotheses could explain such difference between HNS of boar taint and chemical evaluation  
266 of androstenone and skatole. In fact, the first important difference is that HNS measures boar taint while  
267 the present chemical evaluation only quantified androstenone and skatole. Although these two  
268 compounds are the main responsible for boar taint, they are not the only ones (Rius & García-Regueiro,  
269 2001). A general hypothesis which is valid for the observations made in the LS/HA group, is that  
270 regardless of whether androstenone is considered pleasant or unpleasant, its perception is related to the  
271 individual ability and detection threshold of the substance (Font-i-Furnols et al., 2003, 2016; Meier-  
272 Dinkel, Sharifi, et al., 2013). Secondly, some VOCs associated with the HS/MA group, which are absent  
273 in the VOC profiles of LS/HA, might possess unpleasant odor descriptors leading to higher HNS.  
274 Thirdly, interaction between androstenone and skatole (Aluwé et al., 2018; Font-i-Furnols et al., 2003;  
275 Garrido et al., 2016; Mörlein et al., 2016) can also influence the perception of boar taint. Lastly, high  
276 levels of androstenone can produce saturation of the panelist's nose, as experienced by the trained  
277 sensory panel. This temporal lack of perception of androstenone could be confused by inability to  
278 perceive androstenone or non-sensitivity towards this substance. It is further necessary to study the  
279 parameters that influence this saturation that are probably related to the individual conditions of the  
280 panelist, the resting time between samples, the concentration of androstenone of the sample (and maybe  
281 its interaction with other compounds such as skatole) and the matrix used for the evaluation (i.e. fat,  
282 smell strips). Moreover, this could explain why samples in the LS/HA group with androstenone content  
283 greater than 3 µg/g scored lower HNS compared to other samples in this category. However, when  
284 levels of skatole were also high (HS/HA group), it was possible to identify tainted samples by the HNS,  
285 probably due to the less saturation characteristic of skatole, as experienced by the trained sensory  
286 panelists. This might also indicate that probably HS/HA scores are mainly due to high skatole levels.

287 Similarly to what is observed here, Meier-Dinkel et al. (2015) pointed out in their study, that sensory  
288 panelists who were able to clearly identify samples with high skatole content as tainted, had trouble  
289 discriminating some samples with high androstenone from untainted backfat samples.

290 The confrontation of the results obtained for the chemical analysis and that of the sensory evaluation  
291 also shows and confirms a certain level of discordance between both methods. For instance, one sample  
292 with HNS = 1 (*i.e.* weak boar taint) is found in the HS/HA region. Similarly, some fats considered  
293 tainted through sensory evaluation are found in the LS/LA region, although close to the threshold  
294 established.

295 A difference in the type of sample used for the two analyses could in part explain the discordance. In  
296 fact, the chemical analyses were performed on methanolic extracts of liquefied and tissue-free fat, while  
297 sensory analyses were done on native fat (Trautmann et al., 2014). The sensory analyses therefore allow  
298 a good representation of all interactions occurring within the fat, including between fatty acids  
299 constituting the majority of boar fat, and gives a complete vision of the VOCs emitted along with skatole

300 and androstenone (Mathur et al., 2012; Meier-Dinkel et al., 2015; Trautmann et al., 2016; Whittington  
301 et al., 2011). Lastly, the presence of samples considered tainted through sensory evaluation in the LS/LA  
302 region, could be explained by the fact that human olfaction is largely variable with sensitivity ranging  
303 several orders of magnitude between individuals, with individuals having specific anosmia, some  
304 specific hyposmia (reduced olfactory acuity) and others specific hyperosmia (increased olfactory  
305 acuity) (Genva et al., 2019) which affects the obtained results at the detection threshold. Given that  
306 some people present anosmia to androstenone (Font-i-Furnols, 2012), the assessors that carried out the  
307 human nose evaluation were selected to be sensitive to androstenone and skatole, some might even  
308 present specific hyperosmia (detection threshold= 0.2 µg/g for androstenone and 0.05 µg/g for skatole)  
309 and hence grade LS/LA fats as tainted samples.

310 Moreover, only androstenone and skatole have been considered as chemical compounds. Androstenols  
311 and indoles, which are also related to boar taint, were not analyzed and they could help to better explain  
312 HNS scores across the whole set of samples.

313 To better understand part of the disagreement between the human nose evaluation and the chemical  
314 analysis, a deeper look into the fatty acid composition of the samples as well as the emission of VOCs  
315 by the heated fat were performed. In fact, boar fat is a highly complex matrix, firstly due to matrix  
316 effects occurring with the fatty acids impacting the release of boar taint compounds and secondly given  
317 the numerous other VOCs present, which interact and result in a complex smell which cannot be  
318 associated to a unique and constant odor descriptor (Haugen et al., 2012; Trautmann et al., 2014).

### 319 *3.2. Understanding the attributed scores in the human nose evaluation*

#### 320 *3.2.1. Fatty acid composition*

321 Fatty acid composition was analyzed for each fat samples and is summarized in **¡Error! No se  
322 encuentra el origen de la referencia..** For easier interpretation of the results, four categories of HN  
323 appreciation were created to represent the backfat samples: no (HNS = 0), weak (HNS= 0.5 and 1),  
324 moderate (HNS= 1.5 and 2), strong (HNS= 2.5 and 3) boar taint.

325 From this table it can be observed that the fatty acid profiles are made up of a majority of  
326 monounsaturated fatty acids (MUFA) followed by the saturated fatty acids (SFA) and lastly  
327 polyunsaturated fatty acids (PUFA) considerably lower than the two other categories.

328 MUFA and PUFA are predominated by two molecules. In fact, the MUFA make up approximately 46%  
329 of the total fatty acid profile, with C18:1n-9 cis constituting 41.90% to 42.83% of the profile by itself.  
330 Similarly, the PUFA category makes up approximately 16% of the profiles, and is itself made up of  
331 12.43% to 15.83% of C18:2 n-6 cis. For SFA, two molecules are present in much greater quantities

332 compared to the rest of the profile, *i.e.* C16:0 and C18:0 making up approximately 23% and 12.5% of  
333 each group.

334 The overall trends between the three categories and the major constituents of the profiles observed  
335 above have also been found in previous studies on pig carcasses *i.e.* higher SFA and lower PUFA are  
336 related to higher boar taint levels (Liu et al., 2017; Mörlein & Tholen, 2015; Raj et al., 2010).

337 A principal component analysis was then performed exclusively with the fatty acid data of the backfats  
338 to highlight the diversity of fatty acid profiles between samples and link it to the evaluated HNS. From  
339 **¡Error! No se encuentra el origen de la referencia.**, biplot of principal components (PC) 1 and 2, it  
340 appears going from right to left along PC1, that “weak” and “moderate” boar taint samples appear first,  
341 followed by the majority of the “strong” boar taint samples on the left side of principal component 1.  
342 When it comes to the “no” boar taint samples it can be said that these are randomly spread across PC1.

343 The correlation between PC 1 and the variables that compose it was analyzed and the top 10 greatest  
344 contributors to PC 1 were pointed out (represented by arrows in **¡Error! No se encuentra el origen de**  
345 **la referencia.**). PUFAs were the most correlated with positive values for PC 1 and SFA the most  
346 negatively correlated with PC1. When considering tainted samples only (all samples to the exception  
347 of “no” boar taint samples), it seems that the amount of SFA increases along with HNS. On the other  
348 hand, PUFA is inversely correlated to SFA and it can be said that these tend to decrease with increasing  
349 HNS (also visible in **¡Error! No se encuentra el origen de la referencia.**) . What remains unclear  
350 however is the reason behind such spreading along PC1 for the “no” boar taint backfats.

351 To supplement this PCA, the correlation between SFA, MUFA and PUFA and HNS were looked at and  
352 correlations of 0.23, -0.03 and -0.23 were obtained, but were in all cases non-significant. This is in  
353 accordance with the observations made above, however being non-significant they further encourage  
354 the finding of Liu et al. (2017) who states that fatty acids cannot predict the score given during human  
355 nose evaluation.

356 The relationship between fatty acid composition and skatole and androstenone content was also  
357 analyzed (Appendix B). As opposed to previous research, it appears from the data gathered here that  
358 there is no relationship between fatty acids constituting the fat and skatole and androstenone contents.  
359 ANOVAs were performed for each fatty acid in the hope of determining differences based on varying  
360 skatole and androstenone concentrations and to the exception of C8:0, no other fatty acids presented a  
361 significant difference between the groups. When looking at the case of C8:0, it appears that this  
362 molecule is present in very small concentrations (from 0.01 to 0.02% of the profiles), although a  
363 significant difference is observed from a statistical point of view, this does not necessarily mean that  
364 there are biological implications. The authors do not believe in a cause-to-effect relationship between

365 skatole and androstenone concentrations and the variation in C8:0 in the profiles given the low  
366 concentrations and the absence of significance for this molecule in previous studies. Such statement is  
367 supported by the findings of Mörlein & Tholen (2015) who describe C8:0 as a molecule that does not  
368 contribute to discriminating fat samples with varying boar taint levels. Additionally, the correlation  
369 between the fatty acids' classes were determined - the correlations between SFA, MUFA and PUFA  
370 with skatole were 0.12, -0.06 and -0.09, respectively, and with androstenone were -0.08, 0.06 and 0.04,  
371 respectively, all of them non-significant. This weak correlation is explained by the low relation between  
372 the compounds in terms of biosynthesis pathways but also by the low number of samples. In fact, if the  
373 correlation was determined considering all 106 samples initially selected, it would be slightly higher.  
374 In this case, correlations of 0.22, -0.09 and -0.30 are obtained between skatole and SFA, MUFA and  
375 PUFA, respectively and are significant in the case of SFA and PUFA. Correlations of 0.10, 0.20 and -  
376 0.26 are obtained between androstenone and SFA, MUFA and PUFA, respectively and are significant  
377 in the case of MUFA and PUFA.

378 The relationship between fatty acid composition and skatole and androstenone level is a complicated  
379 topic to elucidate. In fact, although several studies have been performed and general trends are similar  
380 between them, it appears that specific relationships observed between fatty acids and boar taint  
381 compounds are not the same from one study to another. Mörlein & Tholen (2015) for example reported  
382 that LS/LA fat samples had higher levels of PUFA compared to HS/HA which hence presented higher  
383 levels of SFA, such as C16:0, C18:0 and high levels of MUFA, such as C18:1. Verplanken et al. (2017),  
384 in their study on the use of rapid evaporative ionization mass spectrometry to discriminate tainted from  
385 untainted carcasses observed similar trends between tainted and untainted boar samples with higher  
386 abundance of some MUFA such as C18:1 and C22:1 in tainted boar fats. On the other hand, Liu et al.  
387 (2017) reported PUFA levels to be positively correlated with androstenone content, while MUFA were  
388 negatively correlated with both androstenone and skatole.

389 From this section it seems that although some trends appear between fatty acids and HNS, these are not  
390 significant. What could explain in part the absence of trends in the above-presented data is that the  
391 backfat samples composing the different taint groups were taken from boars reared under different  
392 production systems with various feeding strategies and different genetics. In fact, as a monogastric  
393 species, the fatty acid composition of pork is a direct reflection of the fatty acid composition in the feed.  
394 Similarly, pigs originated from different breeds, with varying genetics, will have varying fatty acid  
395 composition (De Smet et al., 2004; Johansson et al., 2002; Wood et al., 2008) as well as different  
396 likelihood of presenting boar taint (Xue et al., 1996).

397        *3.3. VOC analysis*

398            *3.3.1. Analysis of skatole and androstenone emissions*

399        The emissions of skatole and androstenone in the headspace of heated fat were related to its associated  
400        content (**¡Error! No se encuentra el origen de la referencia.**). It appears that the emissions in skatole  
401        and androstenone in the headspace of heated fat are a good representation of their corresponding  
402        content. Two other main observations can be made. Firstly, the correlation was stronger for skatole  
403        than androstenone probably because of the higher vapor pressure of skatole compared to androstenone  
404        ( $7.3 \times 10^{-4}$  kPa and  $1.3 \times 10^{-6}$  kPa at 25 °C, respectively). Another explanation could be that some of  
405        the androstenone present in the fat is found as conjugates, less volatile than free androstenone and  
406        subsequently less easily released from the fat matrix (i.e. leading to a poorer correlation). Such  
407        conjugates include androstenone sulfates such as androst-3-enol-3-sulfate and androstenone-4-sulfate  
408        (Bone & Squires, 2021; Squires et al., 2020). Secondly, correlation is lower at higher levels of  
409        androstenone than at lower ones. This could in part explain the difficulty to classify tainted samples  
410        with high androstenone (and low skatole) with the human nose methodology as presented in section  
411        3.1.

412        The correlation between skatole and androstenone content and emissions has already been described in  
413        a previous paper (Burgeon, Markey, et al., 2021) and will not be further developed here.

414        These correlation plots, simply confirm that the skatole and androstenone concentrations perceived by  
415        the human nose during sensory evaluation of boar taint, were overall representative of their actual  
416        content. Part of the explanation between the existing discrepancy between sensory evaluation and  
417        chemical analyses of boar fat must therefore reside elsewhere.

418            *3.3.2. Analysis of general VOC profiles*

419        The best possible representation of the VOC profile of heated fat was performed in this study by  
420        sampling the headspace compounds with a DVB/CAR/PDMS SPME fiber. This fiber is frequently used  
421        to perform untargeted analyses of VOCs given its ability to capture a great variety of VOCs (both in  
422        terms of volatility and polarity). One should however remember that the analyzed VOCs are dependent  
423        of the sampling conditions (amongst which are found the SPME fiber coating) and could therefore differ  
424        from one study to another. In this study, the VOC profiles obtained are composed of 61 molecules  
425        (**¡Error! No se encuentra el origen de la referencia.**). Amongst these are found more than 6 different  
426        chemical families. The profiles are, however, constituted mainly from fatty acids, making up  
427        approximately 73.07% of the VOC profiles, followed by aldehydes, which make up approximately  
428        20.69% of the VOC profiles. Alcohols, alkenes, ketones and furans are the other chemical families  
429        observed.

430 Many of these molecules are known to be typical products of fatty acids oxidation and have already  
431 been previously observed in studies on the VOC profiles of heated fat (Burgeon, Markey, et al., 2021;  
432 Rius et al., 2005; Serra et al., 2014; Sørensen & Engelsen, 2014). In fact, PUFAs such as C18:2n-6 cis,  
433 play an important role in the odor of pork due to the multitude of VOCs that are produced through its  
434 oxidation (Aaslyng & Schäfer, 2008). Typical VOCs produced include alcohols (e.g. pentan-1-ol), fatty  
435 acids (e.g. octanoic acid) and aldehydes (e.g. hexanal) (Domínguez et al., 2019). SFAs on the other  
436 hand, contribute less to the generation of VOCs and hence to the general smell of heated pork. In fact,  
437 these are 100 times less reactive than PUFAs (Parker, 2015). As observed in **¡Error! No se encuentra**  
438 **el origen de la referencia.**, hexadecanoic acid (i.e. C16:0) is found in great abundance in the headspace  
439 of heated fat, which depicts well the smaller reactivity of SFAs. Such free fatty acids, which are not  
440 degraded and found intact in the headspace, have high sensory thresholds which further emphasizes  
441 their low contribution to the smell of heated fat (e.g. C16:0 has detection threshold of 10 000 mg/kg in  
442 oil (van Gemert, 2011)).

443 Although the SPME-GC-MS method used in this study is identical to that used in a Burgeon, Markey,  
444 et al., (2021)'s paper and the results being qualitatively similar, the relative abundance of the  
445 compounds are different. In fact, in this study the VOC profiles are majorly constituted of fatty acids  
446 when aldehydes were the major constituents of the profiles in the above-mentioned study for the  
447 analysis at 150°C.

448 Another difference between the two studies is that skatole is part the VOC profiles obtained here by  
449 analyzing the data in SCAN mode, while this compound was only observed in SIM mode in Burgeon,  
450 Markey, et al., (2021)'s study. This can be explained by the fact that tainted samples with much higher  
451 skatole content were analyzed in this case and is consequently found in higher headspace concentrations  
452 and thus observed in SCAN mode for some samples here. Although the skatole emissions were  
453 significantly correlated to the skatole content as shown in **¡Error! No se encuentra el origen de la**  
454 **referencia.**, it is not surprising that no significant differences exist between the groups presented in  
455 **¡Error! No se encuentra el origen de la referencia.** In fact, the data was gathered here in SCAN  
456 mode and is presented in percentage of the total VOCs profiles where skatole only constitutes minor  
457 percentages.

458 In this article, a focus is brought on the comprehension of which molecules could be responsible of the  
459 varying perception between the different HN appreciation groups. A principal component analysis  
460 (**¡Error! No se encuentra el origen de la referencia.**) was performed here to understand whether  
461 analyzing exclusively general VOC profiles obtained through an untargeted approach can point out  
462 differences between HNS scores.

463 As opposed to the PCA performed with fatty acid content (**¡Error! No se encuentra el origen de la**  
464 **referencia.**), the “no” boar taint backfat samples are less randomly distributed and are found on the

465 right side of PC 1 (to the exception of one sample), whereas the majority of strong boar taint samples  
466 are found on the left side of PC 1. Separation of the other taint groups is less evident. However,  
467 analyzing the top 10 molecules that are the most correlated to PC1 should help to point out trends  
468 between “no” boar taint and “strong boar taint” which have not been perceivable with fatty acid  
469 composition.

470 As it can be noticed by the arrows represented in **¡Error! No se encuentra el origen de la referencia.**,  
471 9 out of 10 top contributors to PC1 are aldehydes, all positively correlated to this principal component.  
472 Most of these molecules have in common that they are direct or indirect products of the oxidation of  
473 fatty acids (Burgeon, Markey, et al., 2021; Serra et al., 2014). For example, characteristic MUFA  
474 oxidation products are found, undec-2-enal and dec-2-enal are C18:1 oxidation products (Domínguez  
475 et al., 2019). Similarly, deca-2,4-dienal is a characteristic VOC of the oxidation of PUFA C18:2.

476 Having the total emitted fatty acids strongly negatively correlated to PC 1, i.e. going towards the  
477 “strong” boar taint samples, joins the idea of having a positive trend between increasing HNS and SFA  
478 content as developed in section 3.2.1 . In fact, as described earlier, SFA being less prone to oxidation  
479 will lead to more intact fatty acids in the headspace of heated backfat samples, as seen here for strong  
480 boar taint samples. Similarly, observing more aldehydes and in particular (*E,Z*)-deca-2,4-dienal  
481 (oxidation product of C18:2) towards the “no” boar taint group joins the idea of increased PUFA content  
482 for lower HNS mentioned in section 3.2.1. These observations which distinguish “no” boar taint  
483 samples from “high” boar taint samples was also confirmed in **¡Error! No se encuentra el origen de**  
484 **la referencia.** In fact, one can notice that the results obtained for the above mentioned molecules are  
485 significantly different between these two HN appreciation groups.

486 To the exception of (*E*)-non-2-enal, which has been attributed various odor descriptors depending on  
487 its concentration, some more pleasant than others – cucumber, green, fat and cardboard odors (Han,  
488 Zhang, & Fauconnier, 2021; Han, Zhang, Fauconnier, et al., 2021; Ross & Smith, 2006; Ullrich &  
489 Grosch, 1987; Zhao et al., 2017), the majority of other aldehydes which appear to have a positive trend  
490 between their emissions and “no” boar taint backfats possess unpleasant odor descriptors. (*E,Z*)-deca-  
491 2,4-dienal, for example, which makes up 7.75% of the total VOC profile has an odor quality described  
492 as “fatty”.

493 One might wonder how are untainted fat samples attributed low HNS (no taint group) given the fatty  
494 odor of most of the molecules described above. The explanation resides in the fact that recognized boar  
495 taint compounds, such as skatole and androstenone, have much greater odor activity values (OAVs)  
496 compared to other molecules cited earlier. In fact, Gerlach et al. (2018) who determined the OAVs of  
497 key molecules in different types of fats, including boar fat found that androstenone and skatole had  
498 OAVs of 25 and 40 respectively as opposed to hepta-2,4-dienal and deca-2,4-dienal which had OAVs  
499 of 2 and 1. This implies that boar taint compounds have a much greater impact on the perception by an



500 assessor evaluating boar taint, compared to oxidation products cited earlier. The results obtained  
501 concerning the correlation between skatole content and HNS and androstenone content and HNS  
502 presented in section 3.1 further supports this idea. Lastly, it should be stated that the appreciation of  
503 fatty odor descriptors is dependent on the food matrix considered. Having fatty and fried odors in  
504 cooked meat is normal and often desired. Hence, these fatty odors most probably do not negatively  
505 contribute to the sensory analysis of backfat.

#### 506 *3.4. Linking content and emissions analysis*

507 Given the observed trends found for both content (fatty acid composition and content in skatole and  
508 androstenone) as well as those observed with VOCs, a PLS-R analysis was used to develop a model  
509 taking into account all analyses performed on the fat matrix to determine whether taken altogether, the  
510 measurements were good predictors of the HNS obtained during sensory evaluation.

511 The PLS regression chosen was one containing only the intercepts and 1 component given that this  
512 yielded the lowest cross-validated root mean square of prediction (RSMEP= 0.8957) and highest  
513 coefficient of determination ( $R^2=0.35$ ). This coefficient of determination indicates that the model has a  
514 certain power of prediction but however is not very strong (an  $R^2$  of 1 signifies that the observed values  
515 can be predicted with 100% accuracy by the model).

516 The VIP scores (VIP= variable importance in the projection) have then been analyzed. The VIP score  
517 is a measure of the contribution of a variable in the model considering the variance explained by each  
518 component (the given variable having a certain impact on each PLS component). It is generally accepted  
519 that a variable should be selected when  $VIP > 1$  (Mehmood et al., 2012). When generating the VIPs, it  
520 was observed that the highest VIP was no greater than 0.15, this indicates that no specific molecule  
521 stands out in the explanation of the model but that the predictive ability of the model is a consequence  
522 of several molecules having a small impact individually. As mentioned in section 3.1, boar taint  
523 perception can be influenced by interactions between skatole and androstenone. This is also true for  
524 interactions with other molecules. However, PLS regression is an additive linear model, interactions  
525 between molecules are therefore not looked at.

526 Yet, the predictive ability of the model could be increased by looking at other factors that are not  
527 considered in this study. Molecules with higher OAVs and recognized as contributing to the smell most  
528 probably play an important role and could explain part of the incoherence between chemical analyses  
529 and sensory evaluation.

530 One must remember that the goal of boar taint detection methods in slaughterhouses is to ensure that  
531 no tainted meat reaches the consumer. To meet this objective, the exact knowledge of the concentration  
532 in boar taint compounds is not required. In fact, what prevails is the overall appreciation of the smell of  
533 pork meat during cooking and consumption. The method that mimics the most this practice is sensory

534 analysis. Additionally, this method is the only one able to perceive all the generated molecules (provided  
535 they are above detection threshold), and the interactions that occur between them (as presented in this  
536 section). Sensory analysis will therefore remain one of the preferred methods for slaughterhouse  
537 detection of boar taint. However, it is necessary that the panelist who perform the analysis are well  
538 trained and sensitive to androstenone and skatole to reduce the false positive and false negative scores.  
539 Furthermore, HNS determine boar taint which, as reported here, is not exactly the same as chemical  
540 analysis of specific and impactful boar taint compounds and this could be a limitation. Other methods,  
541 such as Raman spectroscopy and sensor-based methods, should however still be exploited to offer a  
542 larger and complete variety of options for slaughterhouses in the coming years.

## 543 **4. Conclusions**

544 Although the human nose method was an overall good representation of the results obtained with the  
545 chemical analysis, some existing incoherencies can be found. These are mainly due to the greater  
546 influence of skatole on sensory evaluation, especially in fat with intermediate HNS. Additionally, this  
547 could also be due to the fact that human olfaction is variable.

548 Even though a relationship can be perceived between skatole and androstenone content and fatty acid  
549 content, this relationship was not evident when HNS and fatty acid content were considered, thus,  
550 indicating that the fatty acid composition is not a good predictor of the score attributed during human  
551 nose evaluation.

552 When looking at VOC emissions when heating fat, significant positive correlations between content  
553 and emissions in skatole and androstenone were obtained. Additionally, backfat samples considered  
554 untainted had greater amounts of aldehydes in the VOC profiles as opposed to highly tainted backfat  
555 samples which were correlated to higher amounts of emitted fatty acids in the VOC profiles. This  
556 observation joined the idea that a positive trend exists between HNS and SFA content. SFA are less  
557 prone to oxidation and therefore lead to more intact fatty acids in the headspace. Lastly, the generated  
558 PLS regression pointed out a positive correlation between the actual and the predicted HNS scores  
559 however the predictive ability of the model remained weak, suggesting that other factors play a role in  
560 sensory evaluation. Investigating VOCs with higher OAVs and recognized as contributing to boar taint  
561 was suggested to increase the coefficient of determination of the PLS regression.

562 Altogether, this supports the idea that elevated HNS attributed to tainted meat is mainly due to recognized  
563 boar taint compounds rather than general modifications of VOC profiles composed of pleasant and  
564 unpleasant odors.

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579 **Appendix**

580 *Appendix A. Black and white*

	Panelist 1	Panelist 2	Panelist 3	<b>General</b>
Sensitivity	0.70	0.86	0.86	0.74
Specificity	0.58	1.00	0.87	0.70
Accuracy	0.68	0.90	0.86	0.74
PPV	0.98	0.70	0.96	0.99
NPV	0.54	0.46	0.50	0.51

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588 Appendix B. Fatty acid composition (% , mean±standard error) according to the taint groups based on  
 589 the chemical concentrations of skatole and androstenone: high skatole/high androstenone (HS/HA), low  
 590 skatole/high androstenone (LS/HA), low skatole/low androstenone (LS/LA), high skatole/medium  
 591 androstenone (HS/MA). Means with a different superscript are significantly different (p<0.05)  
 592 according to Tukey-Kramer's test.

593

	HS/HA	LS/HA	LS/LA	HS/MA
C8:0	0.02±0.00 <sup>a,b</sup>	0.02±0.00 <sup>a</sup>	0.01±0.00 <sup>b</sup>	0.01±0.00 <sup>a,b</sup>
C10:0	0.07±0.01	0.07±0.01	0.07±0.01	0.06±0.00
C12:0	0.09±0.01	0.08±0.01	0.08±0.01	0.09±0.01
C14:0	1.24±0.06	1.18±0.05	1.17±0.03	1.25±0.04
C15:0	0.06±0.01	0.08±0.01	0.08±0.0	0.08±0.01
C16:0	23±0.66	22.44±0.35	22.96±0.49	23.96±0.35
C17:0	0.34±0.03	0.41±0.05	0.43±0.06	0.38±0.05
C18:0	13.06±0.47	11.67±0.63	12.37±0.73	13.17±0.67
C20:0	0.23±0.01	0.17±0.02	0.18±0.01	0.20±0.02
C21:0	0.01±0.00	0.02±0.00	0.02±0.00	0.01±0.00
C22:0	0.12±0.01	0.13±0.01	0.12±0.01	0.12±0.02
Σ SFA	38.23±1.04	36.26±0.84	37.48±1.17	39.35±0.97
C14:1	0.02±0.00	0.03±0.00	0.03±0.01	0.04±0.02
C16:1	2.32±0.10	2.53±0.11	2.47±0.18	2.53±0.13
C17:1	0.28±0.02	0.35±0.03	0.33±0.03	0.31±0.04
C18:1 n – 9 cis	42.63±0.78	42.84±0.53	42.65±0.93	42.19±0.59
C18:1 n – 9 trans	0.24±0.01	0.24±0.01	0.26±0.02	0.25±0.02
C20:1 n – 9	0.85±0.04	0.78±0.04	0.77±0.06	0.74±0.03
Σ MUFA	46.34±0.84	46.78±0.57	46.51±1.06	46.06±0.65
C18:2 n – 6 cis	13.51±0.94	14.91±1.01	14.06±0.83	12.85±0.64
C18:2 n – 6 trans	0.11±0.01	0.12±0.01	0.13±0.02	0.14±0.01
C18:3 trans	0.03±0.00	0.03±0.00	0.03±0.01	0.02±0.00
C18:3 n3	0.70±0.06	0.74±0.06	0.69±0.05	0.64±0.04
C20:2	0.62±0.05	0.62±0.01	0.59±0.03	0.51±0.03
C20:3 n – 6	0.10±0.01	0.10±0.00	0.09±0.01	0.09±0.01
C20:4 n – 6	0.26±0.05	0.32±0.04	0.29±0.03	0.26±0.03
C20:5 n – 3	0.01±0.00	0.02±0.00	0.01±0.00	0.01±0.00
C22:5:DPA	0.04±0.01	0.03±0.01	0.05±0.00	0.04±0.00

C22:6DHA:n3	0.06±0.01	0.05±0.00	0.08±0.02	0.03±0.00
Σ PUFA	15.44±1.07	16.96±1.11	16.02±0.94	14.59±0.72
Total	100	100	100	100

594

595 *Appendix C. Mean and standard deviation of the androstenone and skatole contents and human nose scores by class group.*

Class group <sup>1</sup>	HS/HA	HS/MA	LS/HA	LS/LA
n	8	7	7	8
Androstenone (µg/g liquid fat)	4.72±1.68	1.35±0.19	4.11±1.45	0.39±0.35
Skatole (µg/g liquid fat)	0.52±0.21	0.27±0.08	0.06±0.02	0.06±0.06
Human nose score <sup>2</sup>	2.38±0.69	2.36±0.48	1.29±0.95	0.31±0.70

596 <sup>1</sup>high skatole/high androstenone (HS/HA), low skatole/high androstenone (LS/HA), low skatole/low  
597 androstenone (LS/LA), high skatole/medium androstenone (HS/MA)

598 <sup>2</sup> Scores: 0: no boar taint, 1: weak boar taint, 2: moderate boar taint; 3: strong boar taint

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## 835 **Tables:**

836 *Table 1. Fatty acid composition (% mean ± standard error) classified according to the HN appreciation category (no, weak,*  
837 *moderate and strong boar taint). Means with a different superscript are significantly different ( $p < 0.05$ ) according to Tukey-*  
838 *Kramer's test.*

HN appreciation	% of total fatty acid profile			
	No BT	Weak BT	Moderate BT	Strong BT
(n)	(n=7)	(n=5)	(n=6)	(n=12)
C8:0	0.01±0.00 <sup>b</sup>	0.02±0.00 <sup>a</sup>	0.02±0.00 <sup>a,b</sup>	0.01±0.00 <sup>b</sup>
C10:0	0.07±0.00	0.08±0.00	0.07±0.00	0.05±0.00
C12:0	0.07±0.00 <sup>b</sup>	0.09±0.00 <sup>a,b</sup>	0.11±0.00 <sup>a</sup>	0.08±0.00 <sup>b</sup>
C14:0	1.13±0.02	1.21±0.02	1.23±0.02	1.26±0.01

C15:0	0.07±0.00 <sup>a,b</sup>	0.09±0.00 <sup>a,b</sup>	0.10±0.01 <sup>a</sup>	0.06±0.00 <sup>b</sup>
C16:0	23.06±0.21	22.30±0.11	22.48±0.14	23.77±0.12
C17:0	0.38±0.02	0.44±0.02	0.45±0.03	0.33±0.01
C18:0	12.79±0.32	11.42±0.35	11.50±0.19	13.28±0.11
C20:0	0.19±0.01	0.17±0.01	0.17±0.01	0.23±0.00
C21:0	0.02±0.00	0.02±0.00	0.02±0.00	0.01±0.00
C22:0	0.11±0.00 <sup>b</sup>	0.14±0.00 <sup>a,b</sup>	0.15±0.01 <sup>a</sup>	0.11±0.00 <sup>b</sup>
ΣSFA <sup>1</sup>	37.90±0.49	35.95±0.41	36.29±0.29	39.20±0.21
C14:1	0.03±0.00	0.04±0.04	0.05±0.01	0.02±0.00
C16:1	2.43±0.08	2.58±0.09	2.48±0.06	2.48±0.02
C17:1	0.31±0.01	0.36±0.02	0.33±0.01	0.29±0.01
C18:1 n – 9 cis	42.83±0.22	42.62±0.54	41.90±0.37	42.81±0.14
C18:1 n – 9 trans	0.25±0.01	0.24±0.00	0.25±0.01	0.24±0.00
C20:1 n – 9	0.81±0.02	0.75±0.02	0.73±0.03	0.81±0.01
Σ MUFA <sup>2</sup>	46.66±0.28	46.59±0.60	45.74±0.40	46.65±0.15
C18:2 n – 6 cis	13.56±0.27 <sup>a,b</sup>	15.30±0.29 <sup>a</sup>	15.83±0.41 <sup>a</sup>	12.43±0.15 <sup>b</sup>
C18:2 n – 6 trans	0.13±0.01	0.12±0.00	0.14±0.01	0.11±0.00
C18:3 trans	0.03±0.00	0.03±0.00	0.03±0.00	0.03±0.00
C18:3 n3	0.67±0.02 <sup>a,b</sup>	0.79±0.02 <sup>a</sup>	0.82±0.02 <sup>a</sup>	0.62±0.01 <sup>b</sup>
C20:2	0.60±0.01	0.64±0.02	0.59±0.02	0.55±0.01
C20:3 n – 6	0.09±0.00	0.10±0.02	0.10±0.00	0.09±0.00
C20:4 n – 6	0.25±0.01 <sup>b,c</sup>	0.34±0.01 <sup>a,b</sup>	0.35±0.02 <sup>a</sup>	0.24±0.00 <sup>c</sup>
C20:5 n – 3	0.01±0.00	0.02±0.00	0.01±0.00	0.01±0.00
C22:5:DPA	0.09±0.01	0.11±0.01	0.08±0.01	0.06±0.00
C22:6DHA:n3	0.02±0.00	0.02±0.00	0.02±0.02	0.01±0.00
Σ PUFA <sup>3</sup>	15.44±0.31 <sup>a,b</sup>	17.46±0.32 <sup>a</sup>	17.97±0.46 <sup>a</sup>	14.15±0.16 <sup>b</sup>
TOTAL	100	100	100	100

840 *Table 2. GC-MS<sup>1</sup> results of VOCs<sup>2</sup> (% , mean± standard error) found in the headspace of heated fat, sampled with the DVB/CAR/PDMS fiber and detected in SCAN mode. The VOCs are*  
 841 *classified according to the HN appreciation category (no, weak, moderate and strong boar taint). All reference RIs are issued from the NIST Mass Spectrometry Data Center, to the exception of*  
 842 *skatole's which is its injected standard's RI. Means with a different superscript are significantly different (p<0.05) according to Tukey-Kramer's test.*

	CAS <sup>3</sup> number	Reference RI <sup>4</sup> (VF-5ms <sup>5</sup> )	Literature RI	% of total VOC profile			
				No BT <sup>6</sup> (n=7)	Weak BT (n=5)	Moderate BT (n=6)	Strong BT (n=12)
<b>Alcohols</b>							
pentan-1-ol	71-41-0	771	771	0.47±0.10	0.49±0.11	0.34±0.09	0.25±0.08
heptan-1-ol	111-70-6	971	970	0.18±0.04	0.18±0.04	0.12±0.06	0.10±0.03
oct-1-en-3-ol	3391-86-4	980	979	0.45±0.12	0.41±0.15	0.30±0.11	0.18±0.06
octan-1-ol	111-87-5	1072	1071	0.44±0.09	0.44±0.08	0.30±0.09	0.22±0.07
<i>Total alcohols</i>				1.53±0.29	1.52±0.32	1.06±0.34	0.75±0.23
<b>Aldehydes</b>							
Unknown aldehyde	NA	702	NA	0.02±0.01	0.03±0.01	0.03±0.01	0.01±0.01
pentanal	110-62-3	720	717	0.35±0.10	0.38±0.08	0.16±0.04	0.26±0.10
hexanal	66-25-1	797	799	1.32±0.34	1.18±0.22	0.83±0.22	0.99±0.26
( <i>E</i> )-hex-2-enal	505-57-7	850	850	0.17±0.04	0.08±0.04	0.06±0.02	0.06±0.02
heptanal	111-71-7	900	900	0.47±0.11	0.30±0.16	0.38±0.12	0.26±0.07
( <i>E</i> )-hept-2-enal	18829-55-5	955	955	2.52±0.49	2.03±0.44	1.78±0.40	1.27±0.21
benzaldehyde	100-52-7	959	960	0.53±0.05 <sup>a</sup>	0.15±0.07 <sup>b</sup>	0.04±0.04 <sup>b</sup>	0.16±0.04 <sup>b</sup>
octanal	124-13-0	1002	1002	0.64±0.13	0.53±0.10	0.46±0.10	0.40±0.09
( <i>2E,4E</i> )-hepta-2,4-dienal	5910-85-0	1010	1012	2.65±0.28 <sup>a</sup>	1.56±0.22 <sup>a,b</sup>	0.87±0.29 <sup>b</sup>	1.24±0.22 <sup>b</sup>
5-ethylcyclopentene-1-carbaldehyde	36431-51-3	1030	1035	0.17±0.05	0.12±0.06	0.09±0.04	0.06±0.03

2-phenylacetaldehyde	122-78-1	1043	1042	0.14±0.04	0.07±0.02	0.08±0.02	0.08±0.01
( <i>E</i> )-oct-2-enal	2548-87-0	1058	1058	1.07±0.19	0.83±0.17	0.65±0.15	0.56±0.11
nonanal	124-19-6	1104	1104	2.15±0.38	1.74±0.30	1.48±0.30	1.37±0.22
( <i>E</i> )-non-2-enal	18829-56-6	1160	1160	0.56±0.09 <sup>a</sup>	0.36±0.06 <sup>a,b</sup>	0.29±0.07 <sup>a,b</sup>	0.28±0.07 <sup>b</sup>
3-ethylbenzaldehyde	34246-54-3	1163	1168	0.01±0.01	0.00±0.01	0.00±0.00	0.00±0.00
decanal	112-31-2	1205	1205	0.03±0.01	0.01±0.01	0.01±0.01	0.01±0.01
( <i>2E,4E</i> )-nona-2,4-dienal	5910-87-2	1214	1215	0.28±0.04 <sup>a</sup>	0.14±0.04 <sup>a,b</sup>	0.06±0.04 <sup>b</sup>	0.11±0.03 <sup>b</sup>
( <i>E</i> )-dec-2-enal	3913-81-3	1262	1262	2.47±0.36	1.60±0.22	1.29±0.30	1.59±0.27
( <i>2E,4Z</i> )-deca-2,4-dienal	25152-83-4	1294	1295	12.04±1.27 <sup>a</sup>	7.33±0.84 <sup>a,b</sup>	5.87±1.21 <sup>b</sup>	6.37±1.08 <sup>b</sup>
( <i>E</i> )-undec-2-enal	2463-77-6	1364	1365	3.47±0.53 <sup>a</sup>	1.99±0.32 <sup>a,b</sup>	1.50±0.39 <sup>b</sup>	1.87±0.35 <sup>b</sup>
dodecanal	112-54-9	1408	1409	0.11±0.04 <sup>a</sup>	0.02±0.02 <sup>a,b</sup>	0.00±0.00 <sup>b</sup>	0.00±0.00 <sup>b</sup>
tetradecanal	124-25-4	1612	1611	0.08±0.05	0.01±0.01	0.01±0.01	0.00±0.00
hexadecanal	629-80-1	1816	1815	0.00±0.00	0.00±0.00	0.03±0.03	0.05±0.02
<i>Total aldehydes</i>				31.28±4.04 <sup>a</sup>	20.46±2.97 <sup>a,b</sup>	15.96±3.34 <sup>b</sup>	16.99±2.96 <sup>b</sup>
<b>Alkenes</b>							
heptadec-1-ene	6765-39-5	1678	1673	0.08±0.03 <sup>a</sup>	0.00±0.00 <sup>b</sup>	0.00±0.00 <sup>b</sup>	0.01±0.01 <sup>b</sup>
<i>Total alkenes</i>				0.08	0	0.01	0.02
<b>(Emitted) Fatty acids</b>							
2-methylpentanoic acid	97-61-0	984	983	0.00±0.01	0.00±0.00	0.00±0.01	0.00±0.00
hexanoic acid	142-61-1	1021	1020	0.00±0.00	0.09±0.09	0.35±0.27	0.24±0.14
octanoic acid	124-07-2	1184	1180	0.22±0.08	0.23±0.10	0.19±0.09	0.26±0.11
nonanoic acid	112-05-0	1275	1273	0.45±0.10	0.26±0.07	0.13±0.06	0.24±0.06
decanoic acid	334-48-5	1371	1373	0.60±0.17	0.22±0.14	0.30±0.14	0.30±0.13

dodecanoic acid	143-07-7	1565	1565	0.96±0.11	0.61±0.11	0.59±0.09	0.55±0.12
tetradecanoic acid	544-63-8	1768	1768	6.36±0.60	4.65±0.62	5.14±0.68	6.69±1.46
pentadecanoic acid	1002-84-2	1861	1859	0.15±0.05	0.03±0.03	0.04±0.04	0.05±0.02
( <i>Z</i> )-hexadec-9-enoic acid	373-49-9	1953	1953	10.9±1.42	8.44±2.16	12.23±2.00	10.01±1.57
hexadecanoic acid	57-10-3	1974	1972	17.34±1.73	16.63±0.84	19.50±3.07	22.84±4.09
( <i>Z</i> )-heptadec-10-enoic acid	29743-97-3	2075	2073	0.42±0.09	0.43±0.07	0.57±0.12	0.79±0.26
( <i>E</i> )-octadec-13-enoic acid	693-71-0	2161	2163	22.13±6.13	39.15±1.94	37.38±7.99	33.91±6.28
octadecanoic acid	57-11-4	2174	2175	0.76±0.33	2.11±0.88	0.99±0.67	2.05±0.66
Unknown fatty acid	NA	2286	NA	0.15±0.07	0.21±0.07	0.31±0.09	0.19±0.05
<i>Total emitted fatty acids</i>				60.43±4.76 <sup>b</sup>	73.08±3.61 <sup>a,b</sup>	77.71±4.37 <sup>a,b</sup>	78.12±3.69 <sup>a</sup>
<b>Furans</b>							
2-pentylfuran	3777-69-3	991	990	0.95±0.24	0.97±0.26	0.75±0.17	0.57±0.11
2-[( <i>E</i> )-pent-1-enyl]furan	20992-69-2	1000	1000	0.02±0.01	0.01±0.01	0.01±0.01	0.00±0.01
2-heptylfuran	3777-71-7	1192	1193	0.18±0.06 <sup>a</sup>	0.00±0.00 <sup>b</sup>	0.00±0.00 <sup>b</sup>	0.01±0.01 <sup>b</sup>
<i>Total furans</i>				1.15±0.27	0.97±0.26	0.76±0.18	0.59±0.11
<b>Ketones</b>							
heptan-2-one	110-43-0	889	889	0.00±0.00	0.00±0.00	0.01±0.01	0.00±0.00
( <i>E</i> )-oct-3-en-2-one	1669-44-9	1039	1038	0.03±0.02	0.03±0.02	0.00±0.00	0.00±0.00
( <i>E</i> )-non-3-en-2-one	18402-83-0	1139	1137	0.03±0.02	0.04±0.02	0.00±0.00	0.00±0.00
1-phenylhexan-1-one	942-92-7	1462	1459	0.01±0.00	0.01±0.00	0.01±0.00	0.00±0.00
pentadecan-2-one	2345-28-0	1699	1700	1.72±0.16 <sup>a</sup>	0.80±0.15 <sup>b</sup>	0.80±0.11 <sup>b</sup>	0.81±0.14 <sup>b</sup>
heptadecan-2-one	2922-51-2	1902	1900	0.93±0.16	0.56±0.10	0.67±0.13	0.59±0.11
Total ketones				2.72±0.32 <sup>a</sup>	1.44±0.23 <sup>b</sup>	1.48±0.24 <sup>b</sup>	1.40±0.25 <sup>b</sup>



**Others**

Unknown other A	NA	685	NA	1.71±0.24	1.88±0.33	2.36±0.68	1.56±0.27
Unknown other B	NA	910	NA	0.03±0.02	0.01±0.01	0.03±0.02	0.00±0.00
Unknown other C	NA	945	NA	0.03±0.02	0.01±0.01	0.01±0.01	0.00±0.00
2-pentylpyridine	2294-76-0	1197	1201	0.38±0.05 <sup>a</sup>	0.08±0.06 <sup>b</sup>	0.08±0.04 <sup>b</sup>	0.08±0.03 <sup>b</sup>
Skatole	83-34-1	1389	1389	0.00±0.00	0.00±0.00	0.01±0.01	0.01±0.01
[(9Z,12E)-tetradeca-9,12-dienyl] acetate	30507-70-1	1795	1795	0.23±0.05	0.15±0.06	0.13±0.03	0.10±0.03
delta-Tetradecalactone	2721-22-4	1924	1912	0.12±0.05	0.03±0.03	0.02±0.02	0.09±0.04
gamma-Palmitolactone	730-46-1	2104	2104	0.27±0.09	0.34±0.05	0.38±0.10	0.29±0.06
bis(2-ethylhexyl) hexanedioate	103-23-1	2398	2398	0.05±0.02 <sup>a</sup>	0.00±0.00 <sup>b</sup>	0.00±0.00 <sup>b</sup>	0.00±0.00 <sup>b</sup>
Squalene	111-02-4	2833	2833	0.00±0.00	0.02±0.02	0.02±0.02	0.01±0.01
<i>Total others</i>				2.81±0.30	2.52±0.29	3.04±0.68	2.14±0.29
<b>Total</b>				100	100	100	100

843 <sup>1</sup>Gas chromatography – mass spectrometry (GC-MS), <sup>2</sup>Volatile organic compounds (VOCs), <sup>3</sup>Chemical abstracts service (CAS), <sup>4</sup>Retention index (RI), <sup>5</sup>VF-

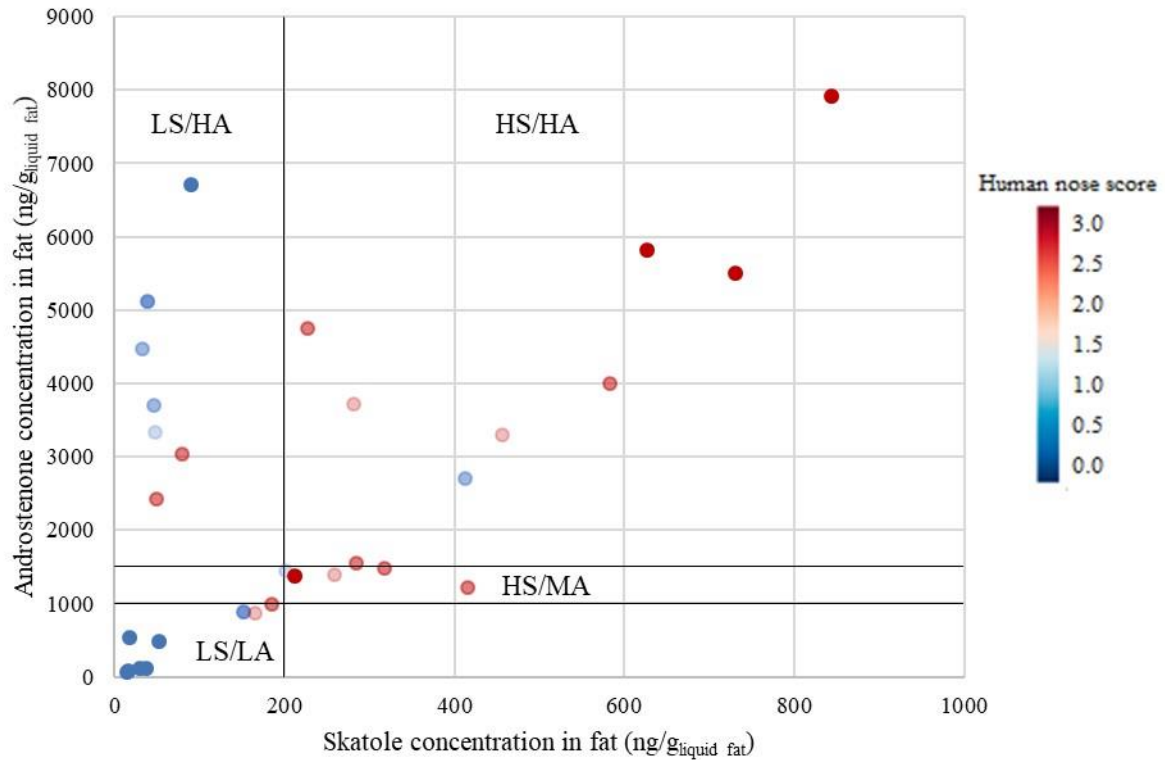
844 5ms is the type of column used, <sup>6</sup>Boar taint (BT)

845

846 **Figure captions:**

847

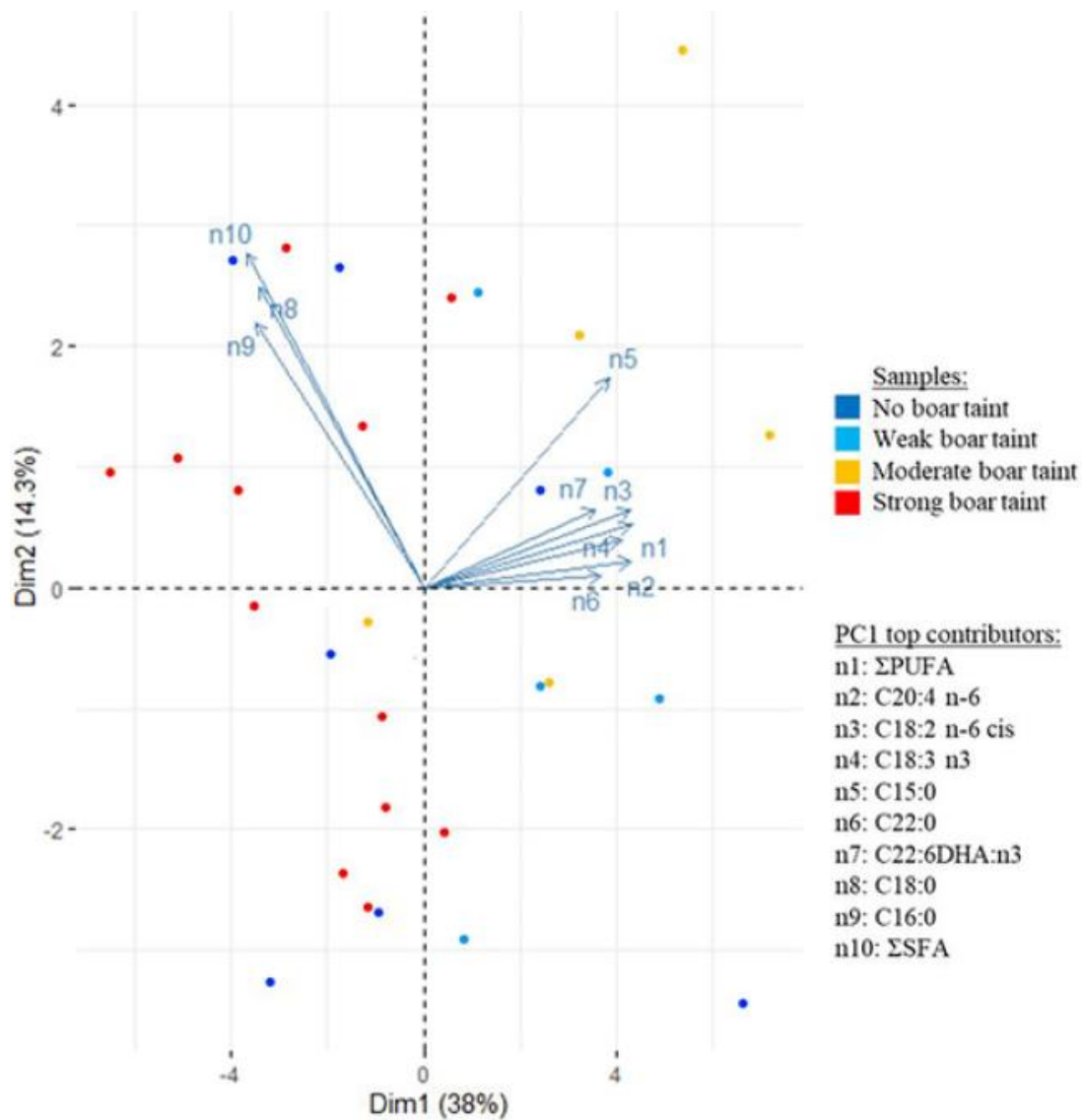
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849

850 Figure 1. Graphical representation combining chemical analysis for skatole and androstenone content  
851 (ng/g liquid fat) on the x- and y-axis respectively, and the score obtained when performing sensory  
852 evaluation through the human nose method. The appreciation corresponding to the human nose score  
853 (HNS) are the following: 0: no boar taint, 1: weak boar taint, 2: moderate boar taint and 3: strong boar  
854 taint perceived for HNS. The black lines allow the separation of the different taint groups – high  
855 skatole/high androstenone (HS/HA), low skatole/high androstenone (LS/HA), low skatole/low  
856 androstenone (LS/LA), high skatole/medium androstenone (HS/MA).

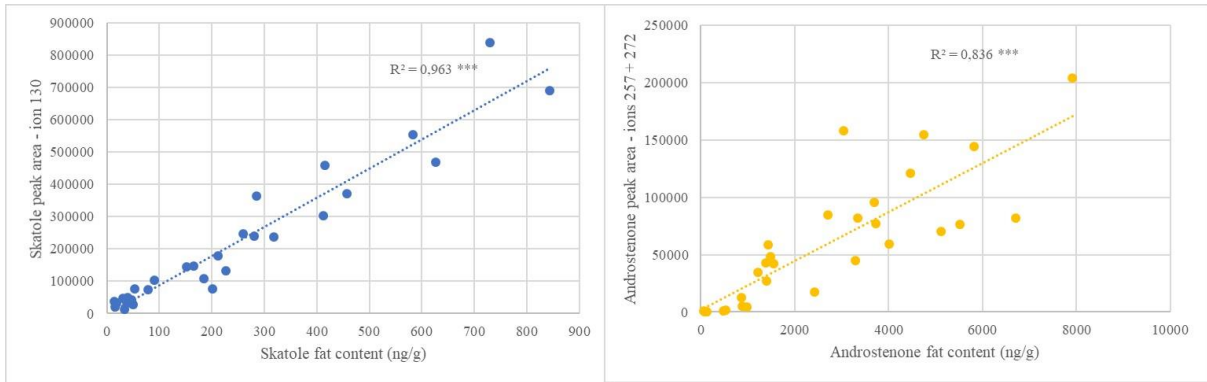
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859 Figure 2. Principal component analysis biplot representing backfats (i.e. the individuals) based on  
 860 their fatty acid composition and the top 10 contributors (i.e. the variables) for principal component 1.  
 861 The backfat samples are represented in four categories based on HN appreciation: no (HNS = 0),  
 862 weak (HNS= 0.5 and 1), moderate (HNS= 1.5 and 2), high (HNS= 2.5 and 3) boar taint. The top  
 863 contributors are represented with arrows (all are very highly significantly correlated to PC1).

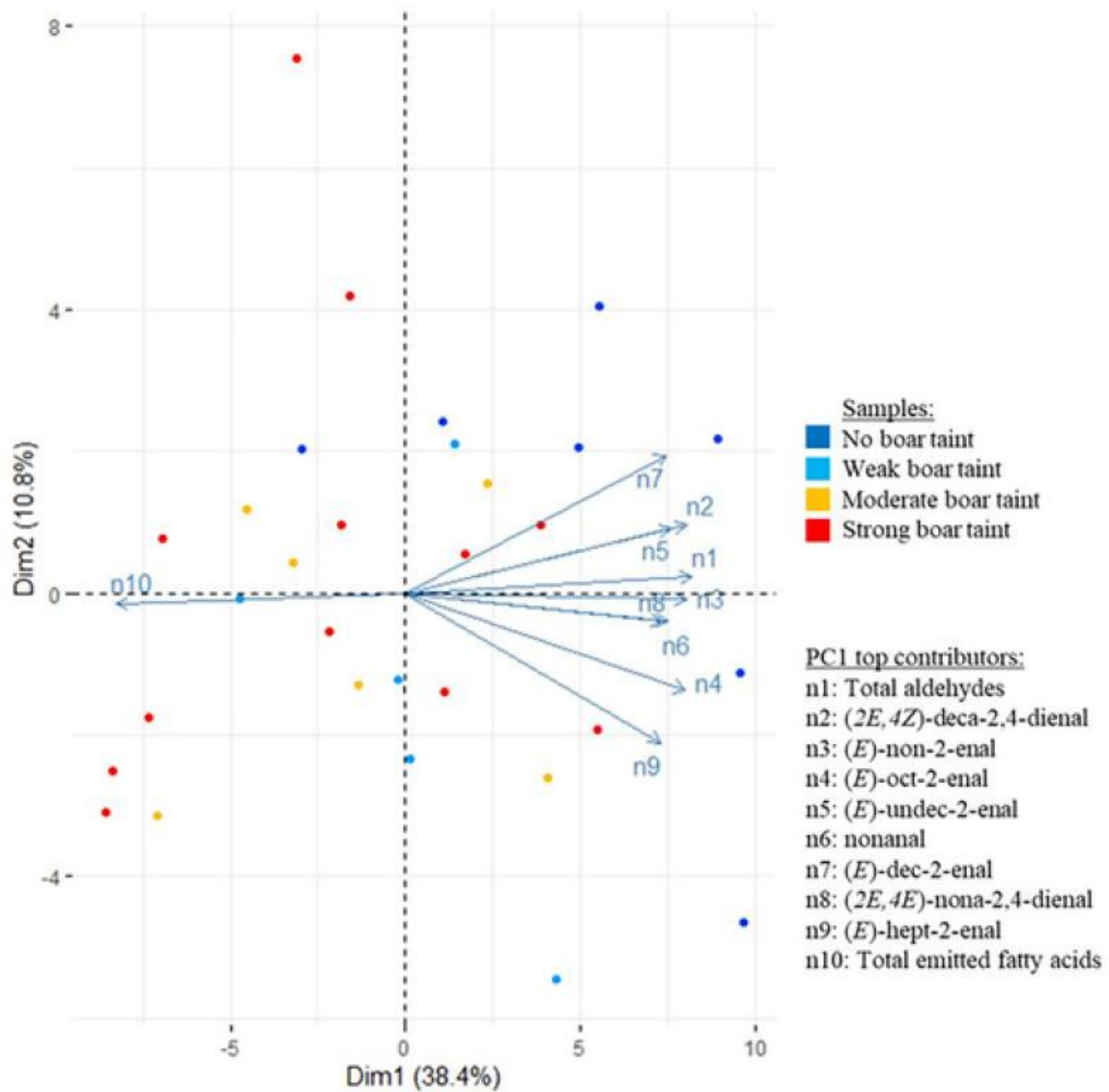
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866 Figure 3. Correlation plots between the emission (peak area) and content (ng/g) of skatole (a) and  
867 androstenone (b) (\*\*\*: P<0.001).

868



869

870 Figure 4. Principal component analysis biplot representing backfats (i.e. the individuals) based on  
 871 their VOC profiles and the top 10 contributors (i.e. the variables) for principal component 1. For  
 872 easier graphical visualization, four categories of HN appreciation were created to represent the  
 873 backfat samples: no (HNS = 0), weak (HNS= 0.5 and 1), moderate (HNS= 1.5 and 2), high (HNS= 2.5  
 874 and 3) boar taint. The top contributors are represented with arrows (all are very highly significantly  
 875 correlated to PC1).