

1 **Isolation of a new taste-active brandy tannin A: structural**  
2 **elucidation, quantitation and sensory assessment**

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17 **Highlights**

- 18 • Original approach to search for taste-active compounds in spirits.
- 19 • Identification of a new taste-active compound: brandy tannin A.
- 20 • Brandy tannin A is an oxidation product of vescalagin.
- 21 • Brandy tannin A is mostly present in cognacs.

22 **ABSTRACT**

23 Enjoying a glass of spirits can be one of the delights of life. While it is well known that their  
24 taste improves during barrel aging, the molecular explanations of this phenomenon remain  
25 largely unknown. The present work aimed at searching for taste-active compounds formed in  
26 spirits during aging. An untargeted metabolomic approach using HRMS was applied on “eau-  
27 de-vie” of cognac. A fractionation protocol was then performed on brandies to isolate a  
28 targeted compound. By using HRMS and NMR, its structure was elucidated for the first time.  
29 This new ellagitannin, called brandy tannin A, considerably increased the sweetness of spirits  
30 at 2 mg/L. After development of an LC-HRMS quantitation method, it was assayed in various  
31 spirits and was detected mainly in cognacs up to 7 mg/L. These findings demonstrate the  
32 sensory contribution of this compound and more generally the relevance of combining  
33 metabolomics and separative techniques to purify new taste-active compounds.

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35 **Key words:** Ellagitannin; sweetness; taste-active compounds; quantitation; oak aging.

## 36 **1. Introduction**

37 Taste is a sense involved in the chemical detection of compounds likely to develop nutritional  
38 or toxic properties. Conventionally, humans can distinguish five basic flavours: salty, sour,  
39 bitter, sweet and umami. Many taste-active molecules, belonging to various chemical  
40 families, have been identified in numerous plants (Kinghorn, 1987). The investigation of these  
41 products in natural matrices such as foods or beverages is a major challenge for chemists.  
42 Such studies are particularly relevant in oenology since they allow a better understanding of  
43 the taste of wines and spirits.

44 In recent years, oenological research has enabled the molecular characterization of non-  
45 volatile compounds involved in the perception of taste, as well as tactile sensations. To isolate  
46 such sensory-active molecules, several strategies have been developed. First, inductive  
47 fractionations guided by sensory analysis were implemented by using various separation  
48 techniques (Frank et al., 2006; Marchal, Waffo-Tégou, et al., 2011). Another strategy was to  
49 search for structural analogues to already known taste-active compounds on the basis of their  
50 putative empirical formulas (Gammacurta et al., 2019; Marchal, Génin, et al., 2015). Then,  
51 analogues were isolated following targeted purification with liquid chromatography-high  
52 resolution mass spectrometry (LC-HRMS) screening and were tasted to determine their  
53 sensory properties. More recently, a new method that combines untargeted and targeted  
54 approaches in the search for new natural products was proposed and applied to spirits  
55 (Winstel et al., 2021). However, the molecular determinants associated with sweet or bitter  
56 taste have only been partially elucidated. These perceptions are linked in particular to the  
57 presence of several compounds released during winemaking by grapes (Cretin et al., 2019;  
58 Fayad et al., 2021), yeasts (Marchal, Marullo, et al., 2011) or by oak wood (Gammacurta et  
59 al., 2019; Marchal, Cretin, et al., 2015; Saucier et al., 2006).

60 Used for a long time as shipping devices, oak barrels are now mainly used for producing  
61 wines and spirits, owing to the physico-chemical and sensory changes they cause. The non-  
62 volatile compounds associated with modifications of colour (Chassaing et al., 2010), tactile  
63 sensations (Glabasnia & Hofmann, 2006) and taste (Marchal, Waffo-Téguo, et al., 2011) that  
64 are consecutive to the aging of wines and spirits have been studied in recent decades.  
65 Empirically, the observations of winemakers suggest an increase in sweetness of wines and  
66 spirits during oak wood aging. This observation has been confirmed (Marchal et al., 2013),  
67 interpreted at the molecular level, and was found to be due to the release of glucosylated and  
68 galloylated triterpenoids called quercotriterpenosides (QTT) (Gammacurta et al., 2019;  
69 Marchal, Waffo-Téguo, et al., 2011). On the contrary, excessive or inappropriate use of oak  
70 wood can increase bitterness in wines and spirits. This phenomenon is mostly attributed to  
71 polyphenols such as lignans (Marchal, Cretin, et al., 2015) and coumarins (Winstel et al.,  
72 2020). Moreover, the ellagitannins of oak wood have been extensively studied for their  
73 possible health effects (Auzanneau et al., 2012; Cardullo et al., 2020; Georgess et al., 2018),  
74 as well as for their sensory properties (Chira et al., 2015).

75 Even if the bitter characteristics of the main ellagitannins have been suggested and their  
76 gustatory detection threshold established (Glabasnia & Hofmann, 2006), the concentrations  
77 observed in wines are significantly below these thresholds. In spirits, the main hydrolysable  
78 tannins, such as vescalagin, castalagin, roburins A to E and grandinin, have been identified  
79 (Gadrat et al., 2020; Puech et al., 1990). However, they seemed to be extracted from oak  
80 wood into “eau-de-vie” from the beginning of barrel aging and were quickly degraded (Viriot  
81 et al., 1993). Likewise and by comparison with the concentrations obtained in various spirits,  
82 no clear correlation could be established between these compounds and the bitterness  
83 sometimes perceived in oaked brandies (Gadrat et al., 2020; Glabasnia & Hofmann, 2006).  
84 Since spirits are a highly complex matrix characterized by an ethanol concentration that is

85 usually between 36% to 55% (v/v), ellagitannins may be involved in various chemical  
86 reactions, including hydrolysis, solvolysis and oxidation. For instance, Quideau *et al.* reported  
87 that in an ethanol solution of vescalagin, the C-1 epimer of castalagin was converted to  $\beta$ -1-*O*-  
88 ethylvescalagin (Quideau *et al.*, 2005). This ethanol adduct has been identified in wines  
89 (Saucier *et al.*, 2006), but never in spirits. Moreover, Fujieda *et al.* discovered oxidation  
90 products of castalagin, named whiskey tannins A and B, in Japanese whiskey (Fujieda *et al.*,  
91 2008). Their structure suggested that they were formed by regioselective oxidation of the  
92 pyrogallol ring linked at the glucose C-1 position of castalagin, and subsequent addition of  
93 ethanol followed by a benzylic acid-type rearrangement. Therefore, spirits aging can lead to  
94 the formation of new chemical species from oak extractables. A recent study provided  
95 confirmation by showing that the diversity of compounds in aged spirits is greater than in the  
96 wood itself (Roullier-Gall *et al.*, 2018). This finding suggests that non-volatiles of oaked  
97 spirits are both native compounds released from wood and molecules newly formed during  
98 aging. Further knowledge is needed to better understand how such changes in the chemical  
99 composition of spirits are linked the overall improvement of their sensory quality during  
100 aging.

101 To search for new taste-active compounds in spirits, the recently proposed combination of  
102 untargeted and targeted approaches was implemented in the present work. First, untargeted  
103 metabolomic profiling by HRMS was carried out on several “eau-de-vie” of cognac of  
104 different vintages. Statistical analyses were performed to evaluate the overall structure of the  
105 metabolomic data and to select compounds potentially newly formed in spirits. Then, a  
106 targeted fractionation protocol, including liquid-liquid extractions, centrifugal partition  
107 chromatography (CPC) and preparative-HPLC, allowed the isolation of new taste-active  
108 compounds that can be further identified, characterized for their sensory properties, and  
109 quantitated in spirits.

## 110 2. Materials and methods

### 111 2.1. Chemicals

112 HPLC grade solvents (acetonitrile, ethanol (EtOH), ethyl acetate (EtOAc), *n*-heptane, propan-  
113 2-ol, methanol and butan-1-ol (BuOH) from VWR International, Pessac, France and methyl  
114 *tert*-butyl ether (MTBE) from Acros Organics, Fisher Scientific, Illkirch, France) and  
115 ultrapure water (Milli-Q purification system, Millipore, France) were used. LC–HRMS  
116 chromatographic separations were performed with deionized ultrapure water, LC–MS grade  
117 acetonitrile and formic acid (Optima, Fisher Chemical, Illkirch).

### 118 2.2. Samples

119 A commercial spirit (Cognac XO) and an oak wood extract (100 g/L), prepared in a hydro-  
120 alcoholic solution (50:50 H<sub>2</sub>O/EtOH) for three days, were screened by LC-HRMS.

121 For targeted compound isolation, a blend of “eau-de-vie” (EDV) of cognac aged in barrels for  
122 19, 20 and 21 years, was used.

123 For quantitative analysis, 36 commercial spirits aged in oak wood were assayed. The second  
124 set of spirits consisted of 9 vintages of EDV of cognac. The samples were not commercial  
125 cognac but EDV still in barrels. They corresponded to a real aging kinetics, i.e. samples of the  
126 same EDV were collected each year in the same barrel. The third set of spirits (Table S1,  
127 Supplementary data) consisted of ten different vintages of EDV of cognac, still in barrels. The  
128 samples came from the same distillery, had undergone similar aging conditions, and had been  
129 matured in used barrels (350 L coarse-grained oak barrels). A sample was collected from five  
130 different barrels for each year (except for 1970 and 1973 for which only four replicates were  
131 available).

132 All spirits were provided by the House of Rémy-Martin. They were diluted with water by a  
133 factor 5 and then filtered at 0.2 µm. This dilution factor was considered when calculating the  
134 final concentration.

135           2.3. *LC-Analysis*

136 For the screening and quantitative analysis, the UHPLC appliance consisted of a Vanquish  
137 system (Thermo Fisher Scientific, Les Ulis, France) with binary pumps, an autosampler and a  
138 heated column compartment. For LC-HRMS analyses and quantitation, a Hypersil Gold C<sub>18</sub>  
139 column (100 × 2.1 mm, 1.9 μm, Thermo Fisher Scientific) was used as the stationary phase,  
140 with water (Eluent A) and acetonitrile (Eluent B), containing both 0.1% of formic acid, as  
141 mobile phases. The flow rate was set at 600 μL/min and the injection volume was 2 μL. The  
142 temperature of the column chamber was set at 30°C in forced air mode. For screening  
143 analysis, eluent B varied as follows: 0 min, 10%; 1.0 min, 10%; 5.0 min, 50%; 5.3 min, 98%;  
144 6.0 min, 98%; 6.15 min, 10%; 7 min, 10%. For the quantitative analysis, eluent B varied as  
145 follows: 0 min, 10%; 1.6 min, 10%; 5.3 min, 35%; 6.1 min, 98%; 7.1 min, 98%; 7.3 min,  
146 10%; 8.3 min, 10%.

147           2.4. *HRMS*

148 For the screening and quantitative analysis, an Exactive Orbitrap mass spectrometer was  
149 equipped with a heated electrospray ionization (HESI-II) probe (both from Thermo Fisher  
150 Scientific). The ionization and spectrometric parameters were designed for each type of  
151 analysis and are summarized in Table 1. Optimization of gas values, voltages and  
152 temperatures applied for ionization and ion transfer was carried out in negative mode.

153 Detection of the targeted compound was based on the theoretical exact mass of its  
154 deprotonated molecular ion ([M - H]<sup>-</sup>) and its retention time. Peak areas were determined by  
155 automatic integration of extracted ion chromatograms (XIC) built in a 3-ppm window around  
156 its exact mass. All data were processed using the Qual Browser and Quan Browser  
157 applications of Xcalibur version 3.0.

158           2.5. *Metabolomic approach*



159 Untargeted metabolomic profiling by HRMS on several EDV of Cognac of different vintages  
160 has already been described in a previous study (Winstel et al., 2021). The analysed samples  
161 corresponded to the third test of spirits (Table S1, Supplementary data). For this approach, a  
162 Q-Exactive Plus mass spectrometer with a HESI-II probe (Thermo Fisher Scientific) was used  
163 and the HPLC appliance consisted of a Waters Acquity I-Class UPLC system (Waters,  
164 Guyancourt, France). Optimization of gas values, voltages and temperatures applied for  
165 ionization and ion transfer was carried out in negative mode (Table 1). After analysing the  
166 spirits samples, Thermo RAW files were exported to the open-source software package  
167 MZmine 2 (2.38 version) for data processing (Pluskal et al., 2010). All the statistical analyses  
168 were carried out using the open-source software R Statistical (Foundation for Statistical  
169 Computing, Vienna, Austria). Results were interpreted by one-way analysis of variance  
170 (ANOVA), using *vintage* as factor. Principal Component Analysis (PCA) was performed on  
171 normally distributed data. K-means clustering was also established for data classification  
172 (package ClassDiscovery). Pearson correlations were then carried out ( $\alpha < 0.05$ , correlation  
173 coefficient  $r > 0.8$ ) and allowed the creation of groups of compounds with a similar evolution  
174 according to the vintages.

## 175 2.6. Extraction and isolation

### 176 2.6.1. Liquid-liquid extractions

177 The blend of EDVs used for isolation was titrated around 64% vol. alc. It was evaporated to  
178 dryness *in vacuo* to remove the ethanol. After evaporation and freeze-drying of four bottles of  
179 750 mL of EDV, a dry extract of 7.35 g was obtained. To start the liquid-liquid extractions,  
180 the extract was first solubilized in 900 mL of milli-Q water. This aqueous extract was washed  
181 twice with 450 mL of *n*-heptane. This aqueous layer was then extracted successively with  
182 MTBE (6×500 mL), EtOAc (5×800 mL) and with water-saturated BuOH (4×800 mL). The  
183 combined organic layers were evaporated *in vacuo*, suspended in water, and freeze-dried

184 twice to obtain brownish powders of MTBE (1.8 g), EtOAc (1.3 g), BuOH (1.2 g) and  
185 aqueous (3 g) pre-purified extracts. They were stored under air- and light-protective  
186 conditions.

### 187 2.6.2. CPC fractionation

188 CPC was performed on a Spot prep II LC coupled with a SCPC-100+1000 (Armen  
189 Instrument, Saint-Avé, France), both controlled by Armen Glider Prep V5.0 software. A 1 L  
190 rotor was used. The solvent was pumped into the column by a 4-way quaternary high-pressure  
191 gradient pump. The samples were introduced into the CPC column *via* an automatic high-  
192 pressure injection valve. All the experiments were conducted at room temperature with UV  
193 detection at 254 and 280 nm. Following the procedure described by Marchal *et al.* (Marchal,  
194 Waffo-Téguo, et al., 2011), the selection of an appropriate biphasic system of solvents was  
195 based on the study of the partition of extract compounds in both phases. Several systems were  
196 tested, and the BuOH extract was fractionated using the ternary biphasic system  
197 EtOAc/propan-2-ol/H<sub>2</sub>O (3:1:3, v/v/v). Separation was carried out by one CPC run of 1.2 g  
198 injection. Experiment was performed at 1200 rpm in ascending mode with a flow rate of 30  
199 mL/min for 135 min for the elution phase and 50 mL/min for 40 min for the extrusion. The  
200 Spot Prep fraction collector was set to 25 mL/min. Every 10 CPC tubes, 200 µL were taken,  
201 evaporated, dissolved in 1 mL of H<sub>2</sub>O/MeOH 90:10 (v/v), filtered and analysed by LC-  
202 HRMS. Ten fractions, named F-I to F-X, were formed according to their similar  
203 chromatographic profile, after being combined, evaporated *in vacuo*, suspended in water, and  
204 freeze-dried.

### 205 2.6.3. Preparative liquid chromatography

206 Preparative HPLC analyses were performed using a Waters Prep 150 LC including a 2545  
207 Quaternary Gradient Module, a 2489 UV/Visible detector (Waters). Final purification of  
208 targeted compound 1 (TC1), which was present in the CPC fractions F-II (83.2 mg), F-III

209 (144.9 mg) and F-IV (62.8 mg), was achieved by preparative HPLC using columns chosen  
210 after LC-HRMS tests. Separations were carried out using a Hypersil Gold C18 (20 mm × 250  
211 mm, 5 μm, Thermo Fisher Scientific) equipped with a Hypersil Gold preparative C18 guard  
212 cartridge (20 × 10 mm, 5 μm, Thermo Fisher Scientific). The mobile phase was a mixture of  
213 ultrapure water (Eluent A) and acetonitrile (Eluent B), both containing 0.1% of formic acid.  
214 The flow rate was set to 20 mL/min. Eluent B varied as follows: 0 min, 10%; 7.4 min, 10%;  
215 44.2 min, 30%; 46.4 min, 98%; 59 min, 98%; 60 min, 10%; 66 min, 10%. Aliquots (20 mg) of  
216 CPC fractions were dissolved in 400 μL of H<sub>2</sub>O/MeOH 60:40 (v/v), 0.2 μm-filtered and  
217 introduced manually into the system. UV detection was performed at 280 nm and  
218 chromatographic peaks were collected manually just after the detector. The pure compound  
219 solution was evaporated *in vacuo* to remove acetonitrile and freeze-dried to obtain a pale-  
220 yellow amorphous powder (10.2 mg).

221 Brandy tannin A (TC1): pale-yellow amorphous powder;  $[\alpha]_D^{25} - 77.8$  ( $c = 0.1$ , MeOH);  
222 HRMS  $m/z$  703.1143  $[M - H]^-$  (C<sub>31</sub>H<sub>27</sub>O<sub>19</sub><sup>-</sup>, 0.6 ppm); <sup>1</sup>H NMR [acetone-D<sub>6</sub>/D<sub>2</sub>O (9:1, v/v),  
223 400 MHz]  $\delta$  1.09 (t,  $J = 7.0$  Hz, 3H), 1.16 (t,  $J = 7.1$  Hz, 3H), 3.47 (dq,  $J = 9.3, 7.0$  Hz, 1H),  
224 3.59 (dq,  $J = 9.4, 7.0$  Hz, 1H), 3.77 (m, 1H), 3.86 (m, 1H), 4.18 (m, 2H), 4.28 (d,  $J = 1.1$  Hz,  
225 1H), 4.40 (dd,  $J = 9.3, 6.7$  Hz, 1H), 4.90 (ddd,  $J = 9.3, 4.3, 2.8$  Hz, 1H), 5.32 (d,  $J = 6.7$  Hz,  
226 1H), 5.40 (s, 1H), 5.48 (d,  $J = 1.0$  Hz, 1H), 6.69 (s, 1H); <sup>13</sup>C NMR [acetone-D<sub>6</sub>/D<sub>2</sub>O (9:1,  
227 v/v), 100 MHz]  $\delta$  201.9, 170.1, 168.6, 168.0, 163.1, 157.9, 146.0, 145.5, 144.5 (2C), 142.6,  
228 136.6, 135.4, 125.4, 125.0, 114.7, 113.5, 111.0, 108.1, 84.6, 80.3, 74.8, 72.7, 67.8 (2C), 65.2,  
229 63.3, 61.9, 45.5, 15.3, 14.0 (Table 2 and Figure S6, supplementary data).

#### 230 2.6.4. NMR experiments

231 NMR experiments were conducted on Bruker Avance II 400 and Avance III 600 NMR  
232 spectrometers equipped with a 5 mm PA BBO and a 5 mm PA BBI probe, respectively. All  
233 1D (proton, carbon, and DEPT-135) and 2D (COSY, HSQC, HMBC, and ROESY) spectra

234 were acquired at 298.15 K in a 9:1 (v/v) acetone-D<sub>6</sub>/D<sub>2</sub>O solvent mixture and were calibrated  
235 using residual undeuterated acetone as an internal reference (<sup>1</sup>H δ 2.05 ppm; <sup>13</sup>C δ 29.8 ppm).  
236 Proton, carbon, DEPT-135, COSY and HSQC spectra were obtained on the Bruker 400 MHz  
237 spectrometer, and HMBC and ROESY spectra were obtained on the Bruker 600 MHz  
238 spectrometer. The following abbreviations were used to describe the multiplicities: s = singlet,  
239 d = doublet, t = triplet, q = quartet, m = multiplet. Data analysis was performed with Mnova  
240 NMR version 14.2.0.

#### 241 *2.7.Method validation for quantitation*

242 A stock solution of brandy tannin A (1 g/L) was prepared in methanol. One range of  
243 calibration was prepared by successive dilutions of this solution in a non-oaked EDV adjusted  
244 to 12% v/v with 0.1% of formic acid, in order to supply calibration samples (1 µg/L, 2 µg/L, 5  
245 µg/L, 10 µg/L, 20 µg/L, 50 µg/L, 100 µg/L, 200 µg/L, 500 µg/L, 1 mg/L, 2 mg/L, 5 mg/L and  
246 10 mg/L).

247 The validation method for quantitating brandy tannin A in spirits was performed by studying  
248 linearity, sensitivity, specificity, intraday repeatability, and trueness. The LC-HRMS method  
249 sensitivity was established using the approach described by De Paepe *et al.* (De Paepe *et al.*,  
250 2013). Limit of detection (LOD) of a molecule is defined as the lowest concentration at which  
251 a reliable and reproducible signal is observed. The signal must be different from a blank  
252 performed under the same conditions. The lowest levels of the calibration curve (from 1 to 20  
253 µg/L) were injected into five replicates. Limit of quantitation (LOQ) is defined as the lowest  
254 concentration of the molecule that can be quantitatively determined by the method, with a  
255 precision lower than 10% and an accuracy (recovery of back-calculated concentrations)  
256 higher than 90%. The working range was based on the LOQ previously determined. A  
257 calibration curve was determined by plotting the areas for each concentration level versus the  
258 nominal concentration. Quadratic regression was used with a 1/x statistical weight. Linearity

259 was evaluated by correlation coefficient ( $R^2$ ) and by deviations of each back-calculated  
260 standard concentration from the nominal value. To determine intraday precision, five  
261 replicates of three intermediate calibration solutions (10  $\mu\text{g/L}$ , 200  $\mu\text{g/L}$  and 10  $\text{mg/L}$ ) were  
262 injected, and the relative standard deviation (RSD%) was calculated. Trueness was checked  
263 by calculating the recovery ratio (between measured and expected areas) from two samples of  
264 EDV (EDV-1; EDV-2). They were chosen among the analysed samples and were spiked with  
265 calibration solution corresponding to an addition of 20  $\mu\text{g/L}$ , 200  $\mu\text{g/L}$  and 10  $\text{mg/L}$  of brandy  
266 tannin A. Interday repeatability was estimated by injections of the same two samples (10  $\mu\text{g/L}$   
267 and 10  $\text{mg/L}$ ) for five successive days. Specificity was assessed by evaluating the mass  
268 accuracy and retention time repeatability. These parameters were determined concomitantly  
269 with the precision and trueness analysis described above.

## 270 *2.8. Sensory analysis*

271 Taste evaluation was performed in a dedicated room, at room temperature (around 20 °C)  
272 (ISO 8589:2010, 2010) and with INAO normalized glass (ISO 3591:1977, 1977). Pure  
273 compounds were tasted by five experts (four women, one man, aged from 24 to 54 years old)  
274 in wine and spirits tasting, at 2  $\text{mg/L}$  in demineralized water (eau de source de Montagne,  
275 Laqueuille, France), as well as in a non-oaked EDV adjusted to 40% (v/v). Experts described  
276 the gustatory perception (bitterness, sweetness, perception of burning and taste of fat) of the  
277 targeted compound using the vocabulary of spirits tasting and were asked to evaluate the  
278 intensity on a scale from 0 (not detectable) to 5 (strongly detectable). The panelists were  
279 informed of the risks and nature of this study and were asked to give their consent to  
280 participate in the sensory analyses. Even though the compound was purified from EDV of  
281 cognac, the experts were advised to spit out the samples after tasting.

282

## 283 **3. Results and discussion**

### 284 3.1. Untargeted metabolomic analysis of spirits to select relevant compounds

285 An untargeted analysis by HRMS was achieved on a series of EDV of cognac of 10 different  
286 vintages from 2015 to 1970 (Table S1, supplementary data), as described in a previous study  
287 (Winstel et al., 2021). After the U-HPLC-HRMS analysis, the data were processed with the  
288 use of MZmine 2 software. Thanks to these treatments, a peaklist of 42,120 negative ions was  
289 obtained between  $m/z$  100 to 1500, then filtered into a peaklist of 331 ions having an  
290 associated data-dependent MS<sup>2</sup> spectrum. PCA of the data was carried out using the peak  
291 areas of the 331 negative ions highlighted by the ANOVA. The vintage effect was clearly  
292 significant on the first axis and the ANOVA showed that it was significant (p-value <0.05) for  
293 321 compounds out of 331 (97%). The compounds were then divided into different groups  
294 showing a similar trend to evolve according to the vintages, by using k-means clustering  
295 followed by Pearson correlations. Of the 321 compounds detected in negative mode and  
296 significantly influenced by the vintage, 298 were assigned to a group among the four created  
297 (Figure S2, supplementary data). Groups 1, 3 and 4 represented 92% of the compounds,  
298 whose concentrations were significantly influenced by the vintage and were generally more  
299 abundant in older vintages. Group 2 was composed of 24 compounds whose contents  
300 increased during 20 years of aging and then slowly decreased.

301 As in the previous study, the statistical groups (Figure S2, supplementary data) revealed the  
302 presence of a wide diversity of molecules in these spirits. For most of them (274/298), the  
303 contents were higher in aged spirits, while the opposite was observed for less than 8%. Such a  
304 result could be explained by two phenomena: a continuous release of oak native compounds  
305 during aging and/or the neoformation of molecules through chemical reactions involving oak  
306 extractables. The aim of this study was to focus on compounds that were formed during spirits  
307 aging, since they cannot be isolated by focusing on oak wood. Therefore, the peaklist obtained  
308 by the MZmine analysis and the data from the statistical analysis were used to target new

309 natural products and attempt to purify them. The compounds of interest were selected  
310 according to three criteria: a significant abundance of the targeted compound, a strong  
311 increase in concentrations in old vintages, and a large gap in intensity between the  
312 concentration in spirits and in oak wood extracts, which could suggest a neoformation rather  
313 than an extraction.

314 First, the 298 compounds were classified in a table according to their intensity in the EDV of  
315 cognac from 1979 to 2015 (data not shown). Then, they were screened by HRMS in the  
316 analysed spirits and in oak wood extracts. Among all the chromatographic peaks, a compound  
317 combining the previously defined criteria was observed and targeted for the rest of the study:  
318 TC1 with a nominal mass of 703. Its concentration increased until 1995 and then slightly  
319 decreased until 1970, while remaining abundant in the EDV of cognac. It was one of the 24  
320 compounds present in group 2. XICs were built by targeting the negative ion at  $m/z$  703.1143  
321 in a 3-ppm window around its theoretical  $m/z$ . LC-HRMS screening revealed the presence of  
322 TC1 only in spirits, which could be explained by a possible chemical reaction in the matrix  
323 during aging (Figure S3, supplementary data). Consequently, its purification protocol was  
324 carried out using the EDV of cognac.

### 325 *3.2. Isolation and Identification of TC1 in Spirits*

#### 326 *3.2.1. Purification of TC1 from “eau-de-vie” of cognac*

327 Metabolomic profiling revealed that TC1 was present at higher levels in the EDV aged in  
328 barrels for 20 years (Figure S4, supplementary data). Its purification was then carried out  
329 from a blend of three EDVs aged in barrels for 19, 20 and 21 years, respectively. First, they  
330 were evaporated to dryness to remove ethanol, which could interfere with subsequent  
331 fractionation steps. After freeze-drying, the second step consisted of sequential liquid/liquid  
332 extractions using MTBE, EtOAc and BuOH to obtain pre-purified extracts. TC1 was mainly  
333 present in the BuOH extract, so this fraction was selected to continue the fractionation. The

334 resulting extract had a complex chromatographic profile with various peaks and co-elutions.  
335 The use of the CPC was necessary to fractionate it and obtain a fraction enriched in  
336 compound  $m/z$  703. Preliminary tests showed that the ternary solvent system EtOAc/propan-  
337 2-ol/H<sub>2</sub>O (3:1:3, v/v/v) in ascending mode allowed the best partition of the sample. Since  
338 many tubes were collected, fractions were constituted by grouping tubes together on the basis  
339 of their LC-HRMS profiles. After solvent evaporation and freeze-drying, 10 fractions (noted  
340 F-I to F-X) were obtained as powder in variable quantities. Fractions F-II, F-III and F-IV were  
341 richer in TC1, so they were submitted to preparative HPLC with UV detection. A first  
342 injection of 5 mg of each fraction revealed that the chromatograms exhibited a refined profile  
343 with only a few peaks detected both in UV a 280 nm. Therefore, a suitable gradient was  
344 chosen for each fraction and F-II, F-III and F-IV were fractionated by successive injections.  
345 The peak corresponding to TC1 was collected manually just after UV detection for each  
346 fraction to give 10.2 mg of a pale-yellow amorphous powder after acetonitrile removal and  
347 freeze-drying.

### 348 3.2.2. *Structural elucidation of TC1*

349 The resolution, mass accuracy and stability offered by HRMS are particularly useful for the  
350 determination of empirical formulas of unknown natural compounds. The HRMS spectrum of  
351 TC1 exhibited a quasi-molecular  $[M - H]^-$  ion at  $m/z$  703.1143. Given the isotopic ratio  
352 (around 35% abundance) and the experimental mass (with a delta of 0.6 ppm) of the  
353 deprotonated ion, the empirical formula C<sub>31</sub>H<sub>28</sub>O<sub>19</sub> was assigned to TC1. To our knowledge,  
354 no compound with this empirical formula has been described in the literature. To investigate  
355 the nature and the sequence of the functional groups, fragmentation was performed on the  
356 pure molecule by non-resonant activation in the higher collision dissociation (HCD) mode  
357 with collision energy of 35 arbitrary units. The fragmentation of TC1 led to the formation of  
358 many ions (Figure S5, supplementary data). The  $m/z$  657.0731 ion, with the molecular



359 formula of  $C_{29}H_{21}O_{18}^-$ , corresponded to a species formed by the loss of a neutral group  $C_2H_6O$   
360 regarding the  $m/z$  703.1143 ion. This group could correspond to a loss of ethanol. Likewise,  
361 the negative  $m/z$  639.0626 ion of the empirical formula  $C_{29}H_{19}O_{17}^-$  corresponded to a  
362 dehydration regarding the  $m/z$  657.0731 ion. Furthermore, the fragmentation spectrum showed  
363 the presence of a negative ion at  $m/z$  523.0513 and could correspond to a species formed by  
364 the loss of a  $C_6H_{12}O_6$  group from the  $m/z$  703.1143 ion, which is characteristic of a hexose. In  
365 addition, an ion at  $m/z$  169.0134 ( $C_7H_5O_5^-$ ) was observed, which may correspond to a gallic  
366 acid. The spectrum also exhibited an ion at  $m/z$  300.9991 ( $C_{14}H_5O_8^-$ ). This might reveal the  
367 presence of the ellagic acid bislactone in the molecule or a structural unit from which this  
368 bislactone could be derived. In addition, ions at  $m/z$  249.0404 ( $C_{12}H_9O_6^-$ ) and  $m/z$  275.0195  
369 ( $C_{13}H_7O_7^-$ ) were detected. They may be 2,2',3,3',4,4'-hexahydroxybiphenyl and 3,4,8,9,10-  
370 pentahydroxydibenzo[b,d]pyran-6-one, respectively. These latter three ions at  $m/z$  301, 275  
371 and 249 are generally characteristic fragments of the main C-glucosidic ellagitannins, such as  
372 vescalagin and castalagin (Bowers et al., 2018). Therefore, by comparing the fragments  
373 obtained with the data in the literature, TC1 could be a C-glucosidic ellagitannin (Engström et  
374 al., 2015; Jourdes et al., 2011).

375

376 A full characterization by NMR was then carried out to identify the structure of TC1, which  
377 was dissolved (5 mg) in a 9:1 (v/v) acetone- $D_6/D_2O$  solvent mixture (Figure S6,  
378 supplementary data). The  $^1H$  NMR spectrum displayed only one aromatic signal resonating at  
379 6.69 ppm, several signals in the downfield sector of the aliphatic chemical shift range between  
380 about 3.5 and 5.5 ppm, which could be due to resonances of protons attached to sugar-type  
381 oxygenated carbon atoms, and two diagnostic triplets just above 1 ppm, each integrating for  
382 three protons. These two signals suggested the presence of two ethoxy units in the structure of  
383 TC1, resulting from chemical transformations involving the spirit ethanol. The  $^{13}C$  NMR

384 spectrum showed 29 distinct carbon resonances out of the 31 carbon atoms presumably  
385 constituting TC1. The observation of two aliphatic carbon signals resonating at 14.0 and 15.3  
386 ppm was in accordance with the presence of two ethyl groups, a finding further corroborated  
387 by the attribution of three signals to (oxygenated) CH<sub>2</sub> carbon resonances at 61.9, 63.3 and  
388 65.2 ppm in the DEPT-135 spectrum. One of these CH<sub>2</sub> signals could be attributed to the  
389 carbon atom of the primary alcohol function of the glucosidic core of TC1 (*i.e.*, C6, Table 2).  
390 The <sup>13</sup>C NMR spectrum also displayed five signals resonating above 160 ppm, which could be  
391 attributed to four carbonyl carbon atoms of ester functions (163.1, 168.0, 168.6 and 170.1  
392 ppm), and a fifth much further downfield signal (201.9 ppm) to a ketone carbon atom.

393  
394 Our hypothesis concerning the *C*-glucosidic ellagitannin nature of TC1 was then further  
395 challenged by performing standard 2D NMR correlation analyses (*i.e.*, COSY, HSQC,  
396 HMBC). The proton signals of the presumed open-chain glucosidic core were assigned on the  
397 basis of <sup>1</sup>H-<sup>1</sup>H COSY data, showing <sup>3</sup>*J* correlations between H1 and H2 (weak), H3 and H4  
398 (strong), H4 and H5 (strong), H5 and the H6's (strong). The latter two diastereotopic protons  
399 H6a and H6b resonated at 3.77 and 3.86 ppm, whose signals overlapped that of the residual  
400 undeuterated water solvent. The COSY data map also revealed the presence of an ethoxy  
401 group through a correlation between the methyl protons at 1.16 ppm (t, *J* = 7.1 Hz) and  
402 methylene protons at 4.18 ppm, and that of another ethoxy group through a correlation  
403 between the methyl protons at 1.09 ppm (t, *J* = 7.0 Hz) and two signals of similar multiplicity  
404 at 3.47 and 3.59 ppm (dq, *J* = 9.4, 7.0 Hz). These two signals, each integrating for one proton,  
405 indicated that they emanate from diastereotopic methylene protons. The same type of signals  
406 was also observed in the <sup>1</sup>H spectrum of the previously described β-1-*O*-ethylvescalagin  
407 (Quideau et al., 2005). Moreover, no correlation was observed with the proton signal  
408 resonating at 5.40 ppm. The signals of protonated carbon atoms could then be assigned on the

409 basis of  $^1\text{H}$ - $^{13}\text{C}$   $^1J$  HSQC data, which notably indicated that C1 and C4 of the glucosidic core  
410 of TC1 would have the same chemical shift at 67.8 ppm. Finally, the analysis of the  $^1\text{H}$ - $^{13}\text{C}$   
411  $^2J/^3J$  HMBC data map enabled us to determine the most likely structure of TC1. A  $^3J$   
412 correlation between C1 and the methylene H1" protons at 3.47 and 3.59 ppm confirmed the  
413 presence of one of the two ethoxy groups on the open-chain glucosidic core, as in the case of  
414  $\beta$ -1-*O*-ethylvescalagin (Quideau et al., 2005). The corollary  $^3J$  correlation between H1 at 4.28  
415 ppm and the methylene C1" at 65.2 ppm was also observed.

416 The other set of methylene H1'" protons resonating at 4.18 ppm were found to correlate with  
417 the ester carbon atom at 170.1 ppm. Acylation of the hydroxy groups at C2, C3 and C5 of the  
418 glucosidic core of TC1 by galloyl-derived units was evidenced by  $^3J$  correlations between H2,  
419 H3, H5 and the carbonyl CI (163.1 ppm), CII (168.0 ppm), CIII (168.6 ppm), respectively.

420 However, the more upfield shift of the carbonyl CI cast doubt on the galloyl nature of the unit  
421 bearing it. Moreover, several remaining carbon signals resonating at 201.9 ppm (ketonic),  
422 157.9 and 142.6 ppm (olefinic), 84.6 and 45.5 (aliphatic) remained to be assigned. In fact, it is  
423 the aforementioned single proton resonance at 5.40 ppm that was the keystone of this  
424 structural determination, since this H5'<sub>I</sub> proton, which is attached to the aliphatic C5'<sub>I</sub>  
425 resonating at 45.5 ppm (HSQC data), correlated with the ketonic C3'<sub>I</sub> at 201.9 ppm, the  
426 olefinic C1'<sub>I</sub> and C2'<sub>I</sub> at 142.6 and 157.9 ppm, and the tertiary alcoholic C4'<sub>I</sub> at 84.6 ppm. In  
427 addition, a  $^3J$  correlation between H5'<sub>I</sub> and the ester carbonyl at 170.1 ppm was also observed.

428 The proximity of H5'<sub>I</sub> with the galloyl-derived unit II was evidenced by  $^2J$  and  $^3J$  correlations  
429 to C-1'<sub>II</sub>, C-2'<sub>II</sub>, and C-3'<sub>II</sub>. Furthermore, the olefinic C1'<sub>I</sub> showed a  $^3J$  correlation with H1 and  
430 a surprisingly strong  $^4J$  correlation with H2 through the ester linkage, and the olefinic C2'<sub>I</sub>  
431 showed a  $^2J$  correlation with H1.

432 All these correlation data suggested that the unit bearing the upfield ( $\alpha,\beta$ -unsaturated) ester  
433 carbonyl CI resonating at 163.1 ppm was a cyclopentenone moiety. The position of the ketone

434 function was established by observing a  $^3J$  correlation between H1 and the carbonyl C3<sub>1</sub> at  
435 201.9 ppm. The cyclopentenone nature of ring I was confirmed by comparing the chemical  
436 shifts of its protons and carbons, and resonance correlations thereof, with those of the same  
437 moiety in whiskey tannin A (Fujieda et al., 2008). In fact, our TC1 is an analogue of whiskey  
438 tannin A, although it is likely not derived from castalagin but rather from its C1-epimer  
439 vescalagin. The  $^3J$  coupling constant between H1 and H2 has a small value of about 1 Hz,  
440 which implies a dihedral angle close to 90° between these two protons and is hence indicative  
441 of its  $\alpha$ -orientation at C1 (Quideau et al., 2005), whereas the same coupling constant in  
442 whiskey tannin A has a value of 3.0 Hz (Fujieda et al., 2008). Furthermore, the ROESY  
443 through-space correlation data map showed signals between H1 and H2, as well as H3, which  
444 are also consistent with an  $\alpha$ -orientation of H1 (Figure 6, supplementary data). The  
445 configurations of the C4' and C5' centres of the cyclopentenone ring I could not be  
446 unambiguously determined, but ROESY correlations between H5'<sub>1</sub> and the ethoxy protons of  
447 the ester function at C4'<sub>1</sub> suggest that H5'<sub>1</sub> and this ester function are *syn*-oriented to one  
448 another. The absence of through-space correlations between H5'<sub>1</sub> and H1 and/or H2 cannot be  
449 used as a strong argument to confirm the  $\beta$ -orientation of H5'<sub>1</sub>, especially since the correlation  
450 between H5'<sub>1</sub> and the  $\beta$ -oriented H1 was also not observed in the NOESY data of whiskey  
451 tannin A (Fujieda et al., 2008). Altogether, the interpretation of our NMR data and the  
452 comparison with literature data on analogous compounds led us to propose the structure  
453 displayed in Figure 1 and Table 2 for TC1, which we name brandy tannin A in reference to  
454 the matrix in which it was identified for the first time.

455

456 Besides the configuration at C1, the other main difference between Tanaka's whiskey tannin  
457 A and our brandy tannin A is the presence of an ethoxy group at this same C1 centre. The  
458 formation of brandy tannin A likely begins with the installation of this ethoxy group onto a

459 starting vescalagin in the ethanol-rich brandy solution (Figure 1). Such a formation of the  
460 resulting  $\beta$ -1-*O*-ethylvescalagin from vescalagin in ethanol was previously described as a  
461 relatively fast, high-yielding and diastereoselective nucleophilic substitution reaction strictly  
462 occurring with retention of configuration at C1 under standard solvolysis conditions (Quideau  
463 et al., 2005). The second step of its formation is probably the oxidative dehydrogenation of  
464 the galloyl-I group leading to the  $\alpha$ -hydroxy-*ortho*-quinone **A**, which can then be subjected to  
465 the nucleophilic addition of ethanol at its most electrophilic carbonyl group. The resulting  
466 dienolic hemiketal **B** might then tautomerize to produce the enonic hemiketal **C**, which can  
467 then undergo a ring contraction *via* a benzilic acid-type rearrangement that forges the C–C  
468 bond between C3'1 and C4'1. Thus, this transformation gives rise to the formation of a  
469 cyclopentenonic ethyl ester, as previously proposed for the formation of whiskey tannins  
470 (Fujieda et al., 2008). Similar dehydrogenation-mediated contractions of ellagitannin galloyl  
471 rings into cyclopentene rings have also been previously reported (Petit et al., 2013; Tanaka et  
472 al., 1990; Wakamatsu et al., 2020). The proposed vescalagin-derived cyclopentenone is in fact  
473 the  $\beta$ -1-*O*-ethyl ether analogue of whiskey tannin B, hereafter referred to as brandy tannin B  
474 (Figure 1). This compound was not observed during our analyses, even though brandy tannin  
475 A certainly derives from it. In the hydroalcoholic brandy solution, the solvolytic cleavage of  
476 its hexahydroxydiphenoyl (HHDP) unit would slowly lead to the formation of brandy tannin  
477 A (TC1).

### 478 3.2.3. *Gustatory properties of brandy tannin A*

479 Brandy tannin A was then dissolved in water and in a non-oaked EDV at 2 mg/L, and the taste  
480 of each solution was characterized in comparison to the same water/EDV as a reference.  
481 Quercotriterpenoside I was used as a sweetness standard since its sensory properties have  
482 already been characterized (Marchal, Waffo-Téguo, et al., 2011). In water, brandy tannin A  
483 exhibited a slight taste of fat, no sweetness, and no bitterness. On a 0–5 scale representing

484 relative taste of fat and sweetness intensity assessed as a consensus between the five panelists,  
485 brandy tannin A scored 2/5 and 0/5, respectively, and QTT I was assessed as 0/5 and 3/5,  
486 respectively. Brandy tannin A was also dissolved in non-oaked EDV to study its influence on  
487 the taste balance of spirits. The control EDV was scored 0/5 for sweetness, bitterness and taste  
488 of fat, but 5/5 for the perception of burning. As a reference, EDV spiked with QTT I (2 mg/L)  
489 was described as sweeter (4/5) and less burning (2/5). Brandy tannin A also modified the taste  
490 of the EDV by significantly decreasing the perception of burning (1/5) and by significantly  
491 increasing that of sweetness (4/5).

492 The results suggested that brandy tannin A developed a taste of fat at 2 mg/L in water, which  
493 modulated the perception of burning of the EDV of cognac and hence improved its overall  
494 taste balance. Moreover, its taste intensity was close to that of QTT I, whose gustatory  
495 detection threshold is relatively low for non-volatile compounds (i.e. 590 µg/L in wine  
496 (Gammacurta et al., 2019), which is much lower than that of glucose, i.e., 4 g/L (Ribéreau-  
497 Gayon et al., 2017).

498 Koga *et al.* found a positive correlation between the antioxidant activity and the aging time of  
499 commercial whiskeys (Koga et al., 2007). In spirits, longer aging leads to a higher  
500 concentration of phenols, especially ellagic and gallic acids and lyoniresinol (Koga et al.,  
501 2007; Winstel & Marchal, 2019). These compounds play an important role in the taste of  
502 whiskey thanks to ROS (Reactive Oxygen Scavenging) and SOD (Superoxide Dismutase)-  
503 like activities (Koga et al., 2011). However, they have mostly been described as bitter  
504 (Marchal, Cretin, et al., 2015; Purwayanti, 2013), so this could not explain why spirits are  
505 known to improve during oak wood aging. Koga *et al.* also considered that there was a  
506 component of spirits which had ROS activity that offered a comfortable aftertaste rather than  
507 an unpleasant one (Koga et al., 2007). Thus, identification of brandy tannin A could provide a  
508 better understanding of the taste balance of spirits aged for a long time in barrels.

### 3.3. Development of an LC-HRMS method to assay brandy tannin A in spirits

509  
510 From a chemical point of view, spirits are complex matrices with thousands of molecules.  
511 Consequently, specific powerful instruments are required to study their composition. Owing  
512 to its mass measurement accuracy and its wide dynamic range, LC-HRMS appeared to be a  
513 reliable technique to quantify brandy tannin A in spirits. To avoid strong matrix effects,  
514 absolute quantitation was carried out by preparing calibration solutions of brandy tannin A in  
515 a non-oaked EDV adjusted to 12% (v/v) with 0.1% of formic acid. In this study, LOD and  
516 LOQ were established at 1 µg/L and 2 µg/L, respectively. A calibration curve was obtained  
517 with a good correlation coefficient ( $R^2$  of 0.999) for a range from 2 µg/L to 10 mg/L, this  
518 validating the linearity of the method. Moreover, all the samples had concentrations that were  
519 in the working range, which confirmed the relevance of the latter. The recovery of back-  
520 calculated concentrations was higher than 90% at each method calibration level, thus  
521 establishing the accuracy of the method. Intraday repeatability for each concentration was  
522 lower than 4.2%. Interday repeatability was not as good at low concentrations (up to 16% at  
523 10 µg/L) but efficient at 10 mg/L (<5%). To overcome this issue, all calibration solutions  
524 were injected for each quantitative analysis of an unknown sample. Two spirits spiked with  
525 stock solutions were also injected. Recovery ratios ranged from 94 to 105%, which is in  
526 accordance with common specifications (Guidance for Industry, 2018). Consequently, these  
527 results established the repeatability and the trueness of the method applied to spirits. Analysis  
528 of the above samples revealed small variations in retention time (<0.02 min) and a mass  
529 deviation lower than 0.9 ppm at various concentrations, guaranteeing the specificity of the  
530 method. All these results validated the LC-HRMS method to quantitate brandy tannin A in  
531 spirits (Table 3).

### 3.4. Quantification of brandy tannin A in spirits

#### 3.4.1. Evolution of brandy tannin A over 8 years

534 Brandy tannin A was quantitated in samples of EDV of cognac of nine different vintages from  
535 the same distillery (Table S7, Supplementary data). The samples were not commercial cognac  
536 but EDV which have been aged in barrels since 2010. A sample was collected each year from  
537 the same barrel from 2010 to 2018, so the 2011 sample corresponds to one year of aging in  
538 barrels, the one of 2012 to 2 years and so on. Brandy tannin A was detected and quantitated in  
539 all spirits at a concentration of 100 µg/L for the sample aged for 1 year in barrels. This result  
540 suggested that it was formed quite quickly after the beginning of aging. The contents of  
541 brandy tannin A were higher in old vintages, reaching a concentration of 2 mg/L for the oldest  
542 sample which had been aged in barrels for 8 years (Figure 2, A). Long barrel aging appeared  
543 to promote the formation of brandy tannin A. Moreover, its reaction rate appeared to be  
544 proportional to its concentration and could be compared to first-order kinetics.

#### 545 *3.4.2. Content of brandy tannin A in various vintages of same spirits*

546 Brandy tannin A was also assayed in the samples of EDV of cognac previously used for  
547 untargeted metabolomic analysis (Table S8, Supplementary data). The concentrations in  
548 Figure 2 correspond to the mean values of the five replicates for the spirits from 2015 to 1990,  
549 and of four replicates for the last two vintages (Figure 2, B). The measured values ranged  
550 from 0.4 mg/L (2015) to 4.2 mg/L (1995). For each vintage, the coefficient of variation  
551 between the replicates was relatively low (from 12.7% to 36.6%), so the heterogeneity  
552 between barrels was not too high, except for the 2005 vintage (47.2%). The evolution of  
553 concentrations for the vintages from 2015 to 2005 was consistent with that of the first series  
554 of spirits. However, brandy tannin A concentrations seemed to follow a bell-shaped curve;  
555 low in the 2015 sample (0.4 mg/L), maximal in the 1995 sample (4.2 mg/L) and lower in  
556 older vintages (e.g. 0.7 mg/L for the 1970 vintage). These results were consistent with its  
557 relative quantitation (Figure S4, Supplementary data) obtained by the untargeted metabolomic  
558 approach, in which the same trend was observed. Even if this was not a strict kinetic study as



559 in the first series of EDV of cognac, this suggested its degradation with barrel aging.  
560 However, this hypothesis needs to be studied more deeply, since the results could also have  
561 been due to modifications of aging practices in the distillery or to changes in barrel supplies.

562 Moreover, six of the ten vintages had concentrations greater than 2 mg/L. Sensory  
563 studies showed significant taste modifications at this concentration, thus demonstrating its  
564 contribution to the taste balance of these spirits. Spirits are known to improve during oak  
565 wood aging and brandy tannin A might play a key role in modulating their taste balance.

#### 566 *3.4.3. Content of brandy tannin A in various commercial spirits*

567 Thirty-six commercial spirits were also analysed to measure the concentration of brandy  
568 tannin A (Table S9, Supplementary data). It was detected in almost all cognacs at  
569 concentrations ranging from 0.03 to 7.7 mg/L but also in two brandies, two whiskeys and one  
570 rum, in smaller quantities (from 0.01 to 0.4 mg/L) (Figure 3). In addition, two Japanese  
571 whiskeys (W-6 and W-7, Figure 3) were analysed since the whiskey tannins A and B have  
572 already been purified from this kind of spirits (Fujieda et al., 2008). Results showed very low  
573 levels of brandy tannin A (< 13 µg/L) in these spirits. The higher brandy tannin A  
574 concentration in the C-7 sample could be due to the significant addition of “boisé” (aqueous  
575 extract of oak wood chips) to this spirit, which is permitted by law for some brandies. The  
576 differences in concentration between the other spirits could be due to the botanical origin of  
577 the wood used for aging. Bourbons are aged in American oak barrels, while cognacs and  
578 brandies are generally aged in French sessile or pedunculate oak barrels. In addition, this  
579 result did not seem surprising since American oaks are known to have much lower  
580 concentrations of ellagitannins than pedunculate oak (Chatonnet & Dubourdieu, 1988).  
581 Additional studies will be necessary to validate this hypothesis. The influence of cooperage  
582 parameters such as the botanical origin of oak wood on brandy tannin A concentrations could

583 be studied. A better control of this parameter could improve the monitoring of oak wood  
584 aging and its sensory effect.

#### 585 **4. Conclusion**

586 This study focused on discovering new taste-active compounds formed during spirits aging in  
587 barrels. For this purpose, an untargeted metabolomic profiling by HRMS in negative mode  
588 was performed on EDV of cognac from several vintages. After statistical analysis, TC1 was  
589 found to be significantly more abundant in spirits than in oak wood, which could suggest its  
590 neof ormation. After the development of a fractionation protocol, brandy tannin A (*i.e.*, TC1)  
591 was identified and purified from a blend of old EDVs of cognac. To our knowledge, its  
592 identification, its presence in spirits, mostly in cognacs, and its sensory properties have never  
593 been described until now. Moreover, its impact on the spirits taste balance was perceived  
594 more strongly by decreasing the burning perception. By determining its gustatory detection  
595 threshold, it might be possible to establish its influence during aging on the taste balance of  
596 old spirits. It would also be interesting to measure its ROS activity to attest to its comfortable  
597 aftertaste. Its concentrations in several EDV of cognac seemed to follow a bell-shaped curve,  
598 suggesting the competition of two phenomena: its formation from a native oak precursor and  
599 its degradation. In both cases, it will be necessary to clarify the chemical species involved, the  
600 reaction mechanisms and the factors that could influence their evolution. The present findings  
601 illustrate the efficiency of our novel method, which allowed the purification of a new  
602 ellagitannin from highly complex mixtures. In future work, such a strategy could be used to  
603 reveal new sensory-active products in natural matrices.

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608

609 **Competing interest statement**

610 The authors declare no competing financial interest.

611

612 **Author contributions**

613 **Delphine Winstel:** Conceptualization, Methodology, Investigation, Validation, Writing –  
614 Original Draft

615 **Yoan Capello:** Conceptualization, Validation, Writing – Original Draft

616 **Stéphane Quideau:** Conceptualization, Validation, Writing – Review and editing,  
617 Visualization

618 **Axel Marchal:** Conceptualization, Validation, Writing – Review and editing, Supervision,  
619 Funding acquisition

620

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787

788 **Abbreviations**

789 EDV : “eau-de-vie”

790 PCA: Principal component analysis

791 TC1: Targeted Compound 1

792 LOD: limit of detection

793 LOQ: limit of quantitation

794 **Appendix A. Supplementary data**

795 **Table S1:** Features of “eaux-de-vie” of cognac used for untargeted LC-HRMS approach.

796

797 **Figure S2:** Representation of different groups of compounds according to their evolution in  
798 48 “eaux-de-vie” of cognac.

799

800 **Figure S3:** Negative LC-ESI-FTMS XIC of an oaked “eau-de-vie” of cognac (A, on the left),  
801 an oak wood extract (B, on the right) corresponding to a negative ion at  $m/z$  703.1143.

802

803 **Figure S4:** Evolution of TC1 in the “eau-de-vie” of cognac from 2015 to 1970. *Error bars*  
804 represent standard deviation of different replicates.

805

806 **Figure S5:** HRMS spectrum of TC1 (with fragmentation 35 eV).

807

808 **Figure S6:**  $^1\text{H}$ ,  $^{13}\text{C}$ , DEPT-135, COSY, HSQC, HMBC, ROESY NMR spectra and  
809 correlation data map of brandy tannin A (TC1) in acetone- $\text{D}_6/\text{D}_2\text{O}$  (9:1, v/v) at 400 MHz and  
810 600 MHz.

811

812 **Table S7:** Individual concentrations of brandy tannin A in 9 vintages of same spirit. All  
813 concentrations expressed in (mg/L).

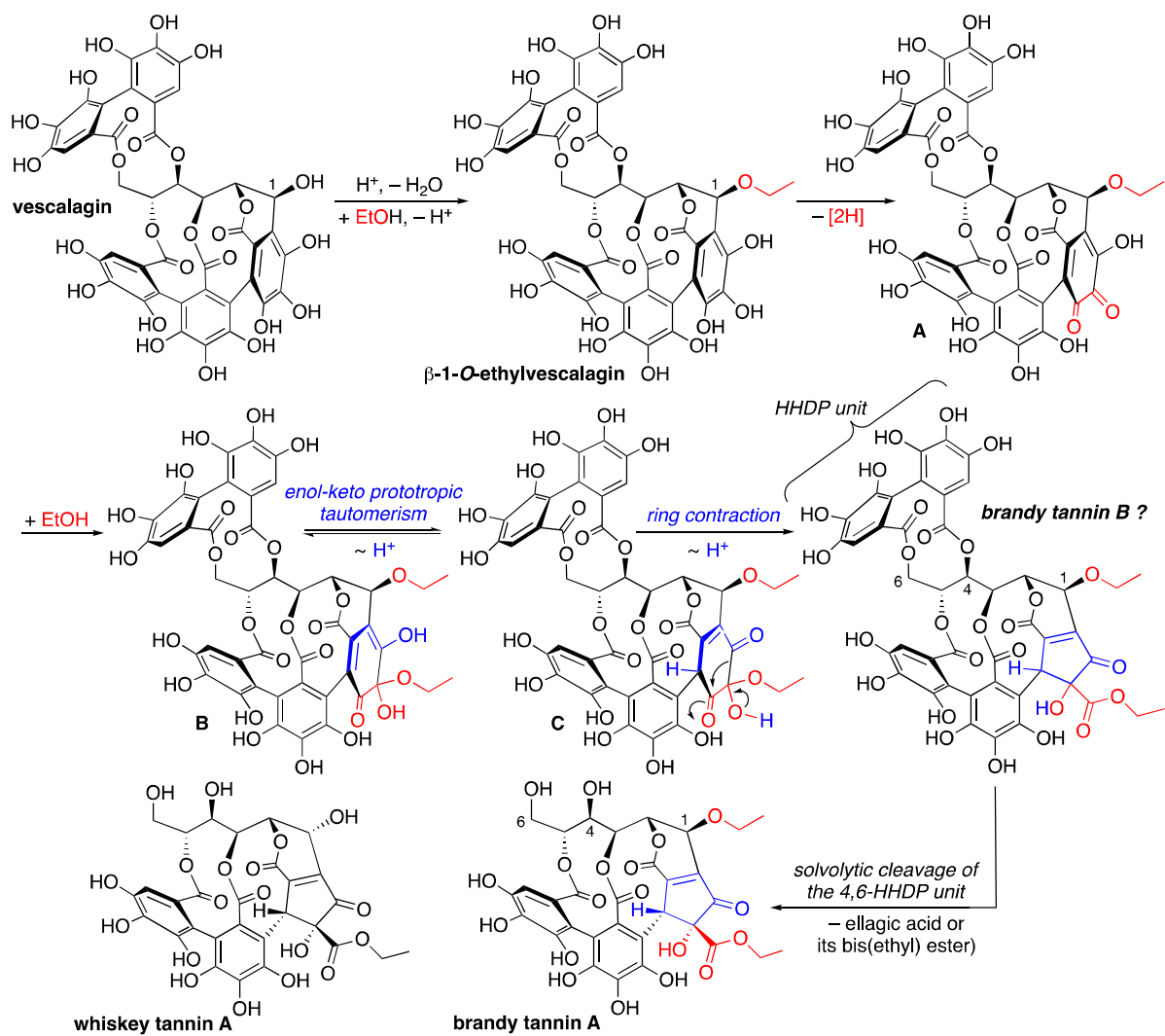
814

815 **Table S8:** Individual concentrations of brandy tannin A in 10 vintages of same spirit. All  
816 concentrations expressed in (mg/L).

817

818 **Table S9:** Individual concentrations of brandy tannin A in 36 commercial spirits. All  
819 concentrations expressed in (mg/L).

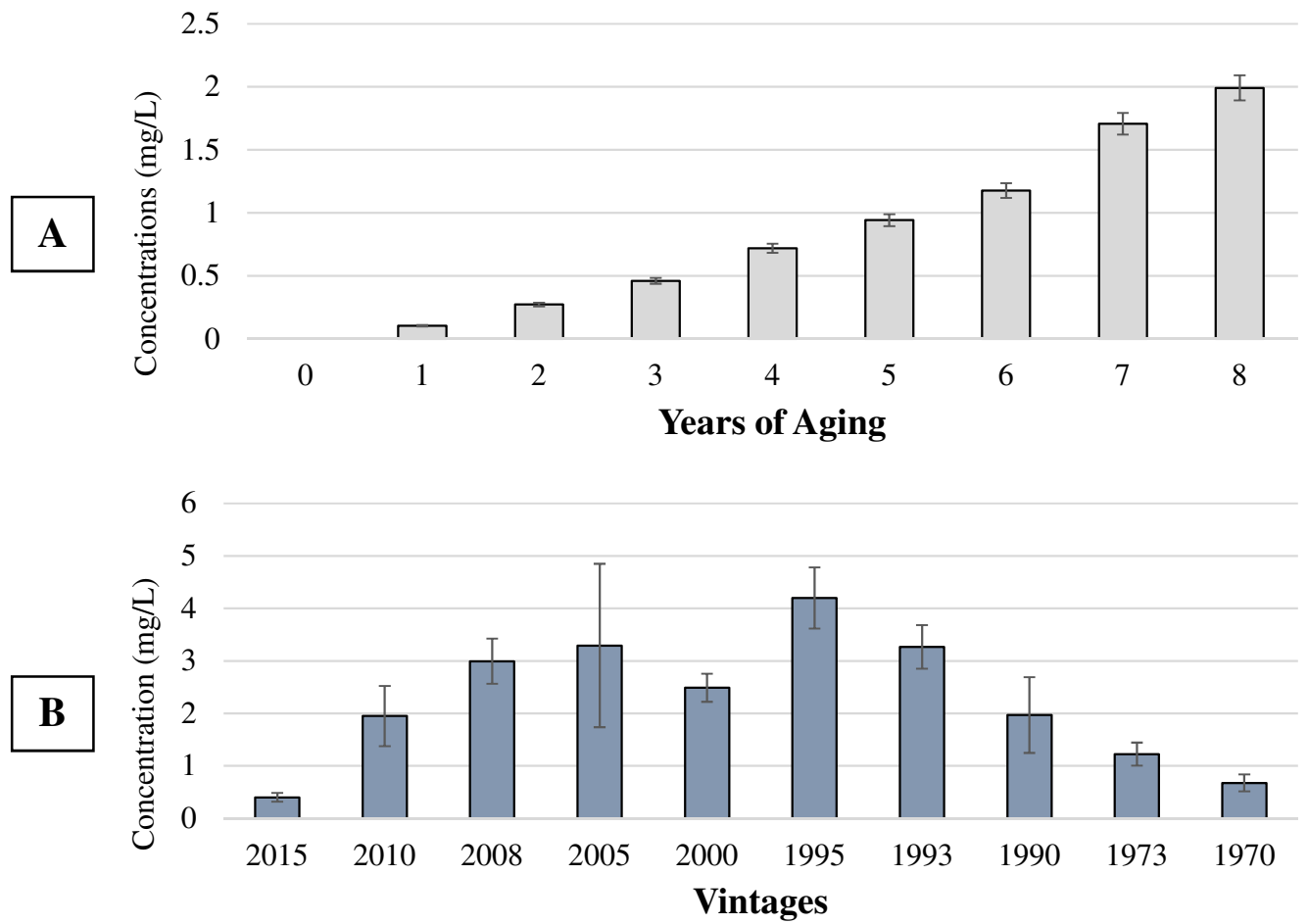
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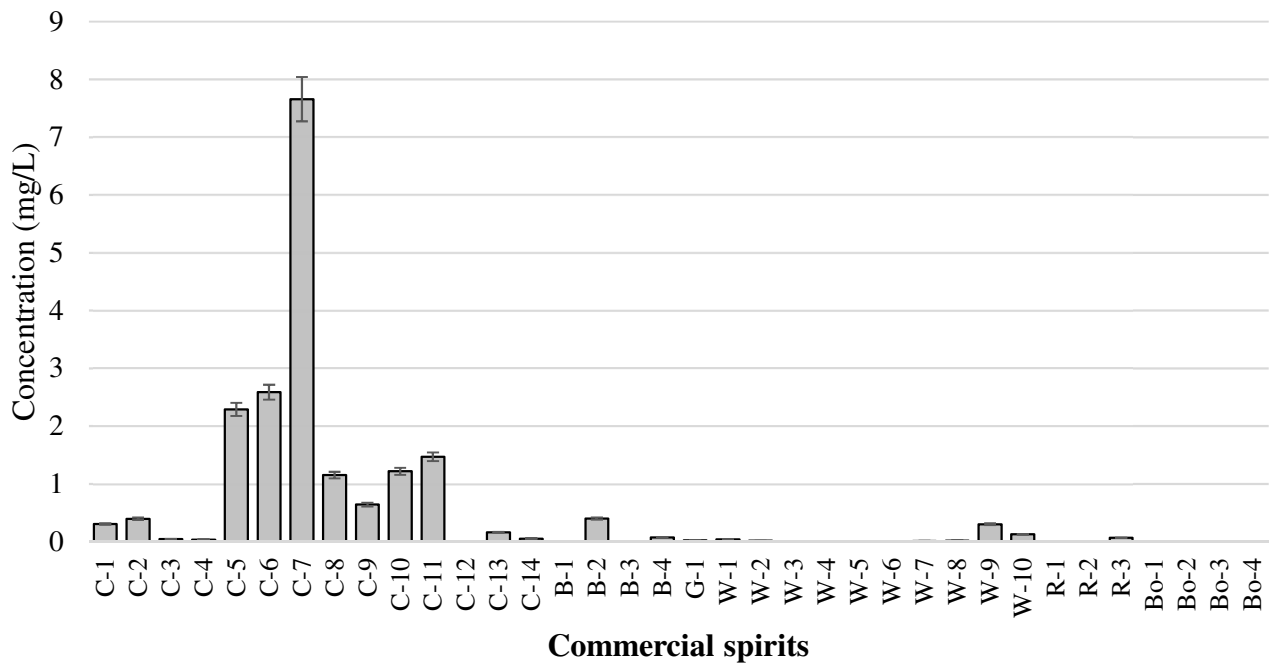
822

823 **Figure 1.** Mechanistic depiction of formation of brandy tannin A from vescalagin.

824



826 **Figure 2:** Concentrations of brandy tannin A over 8 years of aging in barrels (A) and in 10  
 827 vintages of “eau-de-vie” of cognac from same distillery (B).



829

830 **Figure 3:** Concentrations of brandy tannin A in 36 commercial spirits (C: Cognac; B: Brandy;

831 G: Gin; W: Whisky; R: Rum; Bo: Bourbon).

832



833 **Table 1.** Ionization and spectrometric conditions for HRMS analyses.

<b>Ionization mode</b>	<b>Negative</b>			
<b>Mass Spectrometer</b>	<b>Q-Exactive Plus</b>		<b>Exactive</b>	
Use	LC-MS <sup>n</sup> Metabolomic approach		LC-HRMS Screening	LC-HRMS Quantitation
Mass scan	Full MS	dd-MS <sup>2</sup>	Full MS	Full MS
Sheath gas flow <sup>a</sup>	48		70	60
Auxiliary gas flow <sup>a</sup>	11		15	15
Spare gas flow <sup>a</sup>	2		0	0
HESI probe temperature	300 °C		320 °C	350 °C
Capillary temperature	300 °C		350 °C	300 °C
Electrospray voltage	- 3.3 kV		- 3.5 kV	- 3.5 kV
S-lens RF level <sup>b</sup>	55		-	-
Capillary voltage	-		- 25 V	- 95 V
Tube lens voltage offset	-		- 120 V	- 160 V
Skimmer voltage	-		- 20 V	- 18 V
Mass range (in Th)	100 - 1500	200 - 2000	200 - 1000	200 - 1000
Resolution <sup>c</sup>	35,000	17,500	25,000	10,000
AGC value <sup>d</sup>	10 <sup>6</sup> ions	10 <sup>5</sup> ions	10 <sup>6</sup> ions	3.10 <sup>6</sup> ions
Maximum injection time	60 ms	50 ms	-	-
Fragmentation	-	28 eV	-	-

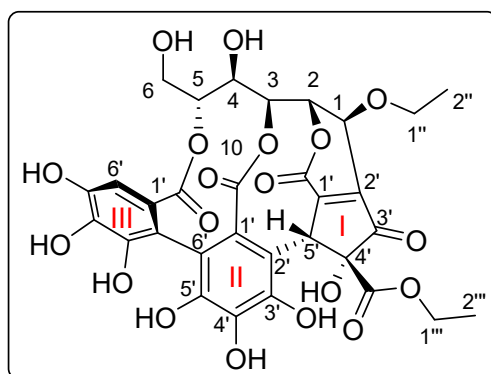
834 <sup>a</sup> Sheath gas, auxiliary gas and spare gas flows (all nitrogen) are expressed in arbitrary units

835 <sup>b</sup> S-lens RF level are expressed in arbitrary units

836 <sup>c</sup> Resolution  $m/\Delta m$ , fwhm at  $m/z$  200 Th

837 <sup>d</sup> Automatic Gain Control

838

839 **Table 2.**  $^1\text{H}$  and  $^{13}\text{C}$  NMR signal assignments of brandy tannin A (TC1).840  
841

Position	$\delta_{\text{H}}$ ( $J = \text{Hz}$ )	$\delta_{\text{C}}$	HSQC	HMBC / COSY
<b>Glucose</b>				
1	4.28 ( <i>d</i> , 1.1)	67.8	C-1	H-2, H-3, C-1'', C-2, C-3, C-1', C-2', C-3'
2	5.48 ( <i>d</i> , 1.0)	80.3	C-2	H-1, H-3, C-1, C-1', C <sub>I</sub> =O
3	5.32 ( <i>d</i> , 6.7)	72.7	C-3	H-2, H-4, C-4, C <sub>II</sub> =O
4	4.40 ( <i>dd</i> , 9.3, 6.7)	67.8	C-4	H-3, H-5, C-6, C-3, C-5
5	4.90 ( <i>ddd</i> , 9.3, 4.3, 2.8)	74.8	C-5	H-3, H-4, H-6, C-4, C <sub>III</sub> =O
6	3.77 ( <i>m</i> ) 3.86 ( <i>m</i> )	61.9	C-6	H-5, C-4
<b>Cyclopentenone</b>				
1' <sub>I</sub>		142.6		H-1, H-2, H-5' <sub>I</sub>
2' <sub>I</sub>		157.9		H-1, H-5' <sub>I</sub>
3' <sub>I</sub>		201.9		H-1, H-5' <sub>I</sub>
4' <sub>I</sub>		84.6		H-5' <sub>I</sub>
5' <sub>I</sub>	5.40 ( <i>s</i> )	45.5	C-5' <sub>I</sub>	C-1' <sub>I</sub> , C-2' <sub>I</sub> , C-3' <sub>I</sub> , C-4' <sub>I</sub> , C-1' <sub>II</sub> , C-2' <sub>II</sub> , C-3' <sub>II</sub> , C=O <sub>Ester</sub>
<b>Aromatics</b>				
1' <sub>II</sub>		125.0		H-5' <sub>I</sub>
1' <sub>III</sub>		125.4		
2' <sub>II</sub>		111.0		H-5' <sub>I</sub>
2' <sub>III</sub>		113.5		H-6' <sub>III</sub>
3' <sub>II</sub>		146.0		H-5' <sub>I</sub>
3' <sub>III</sub>		144.5		H-6' <sub>III</sub>
4' <sub>II</sub>		135.4		
4' <sub>III</sub>		136.6		H-6' <sub>III</sub>
5' <sub>II</sub>		144.5		H-5' <sub>I</sub>
5' <sub>III</sub>		145.5		H-6' <sub>III</sub>
6' <sub>II</sub>		114.7		
6' <sub>III</sub>	6.69 ( <i>s</i> )	108.1	C-6' <sub>III</sub>	C-2' <sub>III</sub> , C-3' <sub>III</sub> , C-

<b>Carbonyls</b>				
C=O <sub>ester</sub>		170.1		H-1'', H-5' <sub>I</sub>
C <sub>I</sub> =O		163.1		H-2
C <sub>II</sub> =O		168.0		H-3
C <sub>III</sub> =O		168.6		H-5, H-6' <sub>III</sub>
<b>Ethyl ether</b>				
1''	3.47 ( <i>dq</i> , 9.3, 7.0) 3.59 ( <i>dq</i> , 9.4, 7.0)	65.2	C-1''	H-2'', C-1, C-2''
2''	1.09 ( <i>t</i> , 7.0)	15.3	C-2''	H-1'', C-1''
<b>Ethyl ester</b>				
1'''	4.18 ( <i>m</i> )	63.3	C-1'''	H-2''', C-2''', C=O <sub>ester</sub>
2'''	1.16 ( <i>t</i> , 7.1)	14.0	C-2'''	H-1''', C-1'''

842

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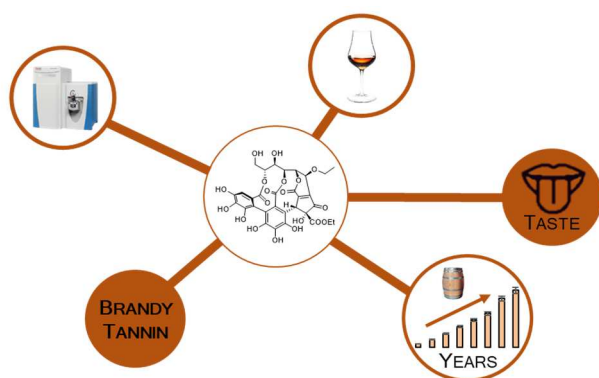
844 **Table 3:** Validation parameters for HRMS quantitation of brandy tannin A in spirits.  
 845

Parameters	Matrix - Spirits		
Sensitivity	LOD ( $\mu\text{g/L}$ )	LOQ ( $\mu\text{g/L}$ )	
	1	2	
Linearity and accuracy	Working range	$R^2$	
	2 $\mu\text{g/L}$ - 10 $\text{mg/L}$	0.999	
Specificity	$t_R$ variation	Mass accuracy	
	0.02 min	0.9 ppm	
	<b>Intraday repeatability</b>		
	<i>10 <math>\mu\text{g/L}</math></i>	<i>200 <math>\mu\text{g/L}</math></i>	<i>10 <math>\text{mg/L}</math></i>
	4.2%	2.7%	2.4%
	<b>Interday repeatability</b>		
Repeatability and trueness	<i>10 <math>\mu\text{g/L}</math></i>	<i>10 <math>\text{mg/L}</math></i>	
	16.3%	4.2%	
	<b>Recovery</b>		
	<i>20 <math>\mu\text{g/L}</math></i>	<i>200 <math>\mu\text{g/L}</math></i>	<i>10 <math>\text{mg/L}</math></i>
	<b>EDV-1</b>	102%	94%
	<b>EDV-2</b>	105%	100%
		102%	94%

846

847

848 **Graphical abstract**



849