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Abstract: Balsamic vinegar of Modena (BVM) and traditional balsamic vinegar of Modena (TBVM) are highly appreciated typical Italian products. The quality control and authentication assurance of both these balsamic vinegars are very important topics. In the recent years, the interest to develop new and standardized analytical procedures, able to further enhance the quality and commercial value of these typical and unique products and to preserve them from possible sophistications and adulterations, is increased. In this work, 76 samples of both BVM and TBVM were analyzed by 1H NMR spectroscopy coupled with multivariate data analysis. The spectral data were analyzed by principal component analysis (PCA), general discriminant analysis (GDA) and classification tree analysis (CTA). The best and very promising model was obtained by a GDA which shows 98.6% of total variance explained by the first canonical function and a predictive capacity of 98.4% with a good separation between clusters. The signals of 5-HMF, α -glucopyranose, malic acid, succinic and acetic acids and a signal at 3.3 ppm referred to the glucose and fructose region were found to be the most statistically significant variables.

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DIPARTIMENTO DI SCIENZE DELLA VITA

Modena, 4 February , 2013

Dear **Katrin Hecht**

LWT- Food Science and Technology Editor

I enclose the manuscript "**Traditional Balsamic vinegar and Balsamic vinegar of Modena analyzed by Nuclear Magnetic Resonance spectroscopy coupled with multivariate data analysis**" and I would ask you to consider it with a view to publishing it in *LWT- Food Science and Technology*.

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Best regards

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PGI BVM and PDO TBVM samples were analyzed by ^1H NMR.

Spectral data were analyzed by PCA, GDA and CTA.

NMR and statistics seem to be a powerful tool in BVM and TBVM characterization.

A model able to classify the PGI and PDO samples was obtained

1 **Traditional Balsamic vinegar and Balsamic vinegar of Modena analyzed by**
2 **Nuclear Magnetic Resonance spectroscopy coupled with multivariate data**
3 **analysis**

4
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25 **Abstract**

26 Balsamic vinegar of Modena (BVM) and traditional balsamic vinegar of Modena (TBVM) are
27 highly appreciated typical Italian products. The quality control and authentication assurance
28 of both these balsamic vinegars are very important topics. In the recent years, the interest to
29 develop new and standardized analytical procedures, able to further enhance the quality and
30 commercial value of these typical and unique products and to preserve them from possible
31 sophistications and adulterations, is increased. In this work, 76 samples of both BVM and
32 TBVM were analyzed by ¹H NMR spectroscopy coupled with multivariate data analysis. The
33 spectral data were analyzed by principal component analysis (PCA), general discriminant
34 analysis (GDA) and classification tree analysis (CTA). The best and very promising model
35 was obtained by a GDA which shows 98.6% of total variance explained by the first canonical
36 function and a predictive capacity of 98.4% with a good separation between clusters. The
37 signals of 5-HMF, α-glucopyranose, malic acid, succinic and acetic acids and a signal at 3.3
38 ppm referred to the glucose and fructose region were found to be the most statistically
39 significant variables.

40

41 **KEYWORDS:** Balsamic Vinegar of Modena , Traditional Balsamic Vinegar of Modena , ¹H
42 NMR spectroscopy, PCA, GDA.

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50 **1. Introduction**

51 The use of high resolution NMR in food authenticity is a subject of great interest in the
52 scientific world, as long as, in the past few years, a widespread growth has been observed, in
53 particular, in the usability of this technique as a fingerprint analysis tool coupled with
54 multivariate data analysis (Bertelli, Papotti, Bortolotti, Marcazzan & Plessi, 2012; Bertelli et
55 al., 2010; Mannina et al., 2012; Lopez-Rituerto et al., 2012; Koda, Furihata, Wei, Miyakawa
56 & Tanokura, 2012; Krishnan, Kruger & Ratcliffe, 2005). The application of NMR in food
57 characterization and control is nowadays very powerful and outstanding due to several good
58 reasons. This technique is well suited to analyze samples without any manipulation, so that, in
59 most cases, extraction and purification are unnecessary, and it is not a destructive analysis.
60 Owing to these advantages, this technique has attended a great development in food science,
61 mainly concerned with the qualitative interpretation of the NMR spectra. This is particularly
62 true for liquid food, such as wine (Papotti et al., 2012), fruit juice (Clausen, Pedersen,
63 Bertram & Kidmose, 2011), beverages (Lachenmeier et al., 2005; Maes, Monakhova, Kuballa,
64 Reusch & Lachenmeier, 2012) and vinegar (Dell'Oro, Ciambotti & Tsolakis, 2012; Van-Diep
65 et al., 2011; Thomas & Jamin, 2009; Boffo, Tavares, Ferreira & Ferreira, 2009; Caligiani,
66 Acquotti, Palla & Bocchi, 2007; Consonni & Gatti, 2004), which may be directly analyzed,
67 after the addition of a deuterated solvent and an internal standard. Other advantages of NMR
68 are the relatively easy and rapid acquisition of data (few minutes are required to acquire a
69 simple ^1H NMR spectrum), the remarkable selectivity and identification of unknown
70 compounds at a molecular level with high reproducibility and repeatability, the ability to
71 furnish structural and quantitative information on a wide range of chemical species in a single
72 NMR experiment, as long as, it is considered as a fingerprint method for authentication
73 analysis.

74 BVM and TBVM are considered typical and well known Italian products, highly appreciated
75 all over the world. Although BVM and TBVM have some organoleptic characteristics in
76 common, they are different products. TBVM is a Protected Designation of Origin (PDO)
77 product (Reg. CE n. 813/2000 April 17, 2000), owing to its typical production procedure and
78 the well-defined area of origin. Nowadays, local vinegar houses, often founded by small
79 family-run business, produce the TBVM, according to the ancient methods of production,
80 whose origins are to be found in the Modenese traditions. Long fermentation and aging
81 procedures, which require expertise and caution in respect of the maturation state of the
82 product, contribute to develop the unique and unmistakable characteristics that we recognize
83 today in this very valuable product. BVM has recently obtained the registration with Protected
84 Geographical Indication (PGI) status, granted by the European Union (Reg. CE n. 583/2009
85 July 3th, 2009). They are both obtained from the alcoholic and acetic fermentation of cooked
86 and concentrated grape musts, and this is the main characteristic that distinguishes balsamic
87 vinegars from other vinegars, which are generally produced from alcoholic solution. TBVM
88 and BVM mainly differ in the aging process and the production procedures. TBVM is aged in
89 characteristic wooden barrels and may be found on the market in two different products
90 according to the aging process: old (>12 and <25 years) and extra old (>25 years). It is
91 allowed the inoculation of colonies of acetic bacteria, while the use of any extra additive is
92 forbidden. BVM is a cheaper product, with maturation in wooden barrels from two months up
93 to 3 years, and it is allowed to add vinegar obtained by wine acetification (10% v/v minimum)
94 and caramel (2% v/v maximum) for color correction (D.M., December 3th, 1965). From the
95 regulatory point of view, the quality control and authentication assurance of both these
96 balsamic vinegars was tested by means of sensorial analysis and by very simple chemical–
97 physical property determinations, like total acidity, density and dry residual. Here arises the
98 interest to develop new and standardized analytical procedures, able to further enhance the

99 quality and commercial value of these typical and unique products and to preserve them from
100 possible sophistications and adulterations. These new approaches, coupled with chemometric
101 approaches, may provide helpful classification models for authentication and commercial
102 quality characterization.

103 Several analytical studies focused their attention on the balsamic vinegars characterization
104 and, some of them aim to investigate the modifications occurring during aging and the age
105 determination (Cocchi, Lambertini, Mancini, Marchetti & Ulrici, 2002; Plessi, Bertelli &
106 Miglietta, 2006; Plessi, Monzani & Coppini, 1989; Del Signore, Stancher & Calabrese, 2000;
107 Theobald, Muller & Anklam, 1998; Antonelli, Chinnici & Masino, 2004; Chiavaro, Caligiani
108 & Palla, 1998; Cocchi et al., 2006; Gullo, C.; Caggia, De Vero & Giudici, 2006). However,
109 they are often time consuming approaches, not compatible with routine analyses. Among
110 many advantages that the ^1H NMR spectroscopy offers, it may simultaneously determine the
111 different metabolites of vinegar in few minutes, as required for food authenticity and quality
112 control. To our knowledge, only few studies have been carried out on Traditional Balsamic
113 Vinegar of Modena (TBVM) and Balsamic Vinegar of Modena (BVM), regarding their
114 quality evaluation and valorization, using NMR (Cirlini, Caligiani & Palla, 2009; Consonni et
115 al., 2008; Consonni, Cagliani & Rinaldini, 2008; Consonni & Cagliani, 2007). Here a
116 characterization of both BVM and TBVM using high-resolution ^1H NMR spectroscopy
117 coupled with multivariate statistical data analysis is presented. Besides, we aim to build a
118 discriminant model able to characterize TBVM according to the aging process, which is
119 actually the most required information for quality assessment, and, nowadays, no objective
120 analytical techniques have been officially defined. The particular climatic characteristics, the
121 soil and the grape varieties typically grown in Modena strongly contribute to make the TBVM
122 a unique and unmistakable product. These characteristics and the particular production
123 procedures used by the local producers, which carry on the Modenese traditions through the

124 art of cooking musts and drawing and topping up procedures among the wooden casks, make
125 it is difficult to obtain statistical models which are representative of the intrinsic variability
126 and peculiarities of TBVM. In this work, we applied supervised pattern recognition
127 procedures never used before, and a very significant number of samples is available, that is
128 non-obvious, also considering the high price of the samples.

129

130 **2. Materials and methods**

131 *2.1. Materials and sample preparation*

132 A total of 76 samples of both TBVM and BVM have been analyzed. Among them, 23 were
133 extra old (>25 years of aging) TBVM, 17 were old (>12 and <25 years of aging) TBVM and
134 36 were BVM of unknown aging (**Table 1**). All the TBVM and several BVM were provided
135 by local vinegar houses, while the other BVM were purchased on the market. All TBVMs and
136 BVMs are labelled as PDO and PGI products respectively. Samples were prepared into the
137 Wilmad NMR tube (5 mm, Ultra-Imperial grade, 7 in. L, 526-PP, Sigma-Aldrich, Milan, Italy)
138 by dissolving 0.1 g exactly weighed of vinegar into 500 μ L of dimethyl sulphoxide- d_6
139 (DMSO- d_6) (Sigma-Aldrich, Milan, Italy), and 20 μ L of tetramethylsilane (TMS) was added
140 as reference compound. Standard compounds for metabolite assignments were from Sigma-
141 Aldrich (Milan, Italy).

142 *2.2. Physical and chemical determinations*

143 Undiluted samples were used for $^{\circ}$ Brix measures, which were carried out with refractometer.
144 Total acidity was determined by the titration Official Method (Resolution OIV-OENO 52-
145 2000). *R ratio*, which is the rate between $^{\circ}$ Brix and Total Acidity % (p/v), and indicates the
146 balance among sweet and sour tones, correlating in this way the density with the acidity
147 (Gullo & De Vero, 2004; Satriani, 2010), was also calculated for TBVM samples. This

148 parameter has always been used as a tool that, if properly interpreted, allowed to conduct
149 correctly vinegar houses and to identify possible outliers, before performing NMR analysis.

150 2.3. NMR spectroscopy

151 To characterize samples ^1H NMR, ^{13}C NMR, bidimensional ^1H - ^{13}C heteronuclear multiple-
152 bond correlation (HMBC) and ^1H - ^{13}C heteronuclear single quantum coherence (HSQC)
153 spectra were acquired with a Bruker FT-NMR Avance 400 spectrometer (Bruker Biospin
154 GmbH Rheinstetten, Karlsruhe, Germany) operating at 400.13 MHz for ^1H . All of the
155 experiments were performed at 300 K and nonspinning. ^1H NMR data were acquired using
156 the Bruker spin-echo sequence “cpmg1d” (Carr-Purcell-Meiboom-Gill, Bruker Library)
157 without water presaturation. Acquisition parameters were as follows: time domain (number of
158 data points), 32K; dummy scans, 4; acquisition time, 3.4210 s; delay time, 3.0 s; number of
159 scans, 64; spectral width, 4789.27 Hz; fidres, 0.1461. Total acquisition time was 7 min and 46
160 s. The assignments have been carried out on the basis of the ^{13}C NMR,
161 HMBC and HSQC analyses. The acquisition parameters of the ^{13}C NMR experiments were as
162 follows: number of scans, 8K; dummy scans, 4; time domain (number 142 of data points),
163 32K; spectral width, 22075.055 Hz; acquisition time, 0.7422 s; delay time, 1.5 s; fidres,
164 0.6737 Hz. Total acquisition time was 5 h, 14 min, and 59 s. The acquisition parameters of the
165 HMBC experiments were as follows: number of scans, 32; dummy scans, 16; time domain,
166 3K in the acquisition or direct HMBC dimension F2 (^1H) and 100 in indirect HMBC
167 dimension F1 (^{13}C); spectral width, 5592.841 Hz in F2 (^1H) and 20124.465 Hz in F1 (^{13}C);
168 digital resolution, 1.8206 Hz in F2 (^1H) and 201.245 Hz in F1 (^{13}C); acquisition time, 0.2747
169 s; delay time, 0.5 s; HMBC delay time, 62.5 ms. Total acquisition time was 82 min and 11 s.
170 The acquisition parameters of the HSQC experiments were as follows: number of scans, 4;
171 dummy scans, 12; time domain, 1K in the acquisition or direct HSQC dimension F2 (^1H) and
172 256 in indirect HSQC dimension F1 (^{13}C); spectral width, 5995.204 in F2 (^1H) and 19118.721

173 F1 (^{13}C); digital resolution, 5.855 Hz in F2 (^1H) and 74.682 Hz in F1 (^{13}C); acquisition time,
174 0.0854 s; delay time, 1.5 s. Total acquisition time was 27 min and 50 s. The chemical shifts
175 were reported as δ_{H} (ppm) relative to TMS.

176 *2.4. Spectral calculation*

177 The classification models were obtained using the ^1H NMR spectra as intensity. The
178 application of the ^1H NMR technique to vinegar samples generates very complicated spectra
179 that need to be previously processed and subsequently analyzed by chemometric methods.
180 Each spectrum generated a 16K data points corresponding to time domain, that is the number
181 of points acquired and digitalized by the instrument along the spectral width, and then
182 converted into a frequency domain spectrum by Fourier transform; these files were collected
183 in a data set consisting of 16K variables and 76 samples. All ^1H NMR spectra were phased
184 and calibrated using the TMS signal by the XWinNMR software package (Bruker Biospin
185 GmbH Rheinstetten). To reduce the inhomogeneous proton NMR chemical shift, all spectra
186 were aligned using the toolbox IcoShift 1.0 for MATLAB (Mathworks Inc., Natick, MA, USA)
187 (Savorani, Tomasi & Engelsen, 2009). Finally, the spectra were baseline corrected by
188 PLS_Toolbox version 5.2.2 for use with MATLAB (eigenvector Research Inc., Wenatchee,
189 WA, USA). All the spectral regions devoid of signals and the residual solvent ($\text{DMSO-}d_6$)
190 signal (region from 2.45 to 2.55 ppm) were not considered. The resulting data set refers to the
191 complete spectral region (12362 variables). Two other data sets have been prepared, the first
192 one referred to the low-frequency spectral region between 0.65 and 2.70 ppm (2586 variables),
193 which principally contains the signals of acidic and aliphatic compounds, and the second one,
194 which contains the signals of the midfrequency region, between 2.70 and 5.50 ppm (3499
195 variables).

196 *2.5. Statistical analysis*

197 Before the spectral analyses, all data were normalized, mean-centered, and scaled by the
198 pareto-scaling method (Winning et al., 2009). To achieve a reliable classification,
199 unsupervised and supervised pattern recognition procedures were applied to the data sets.
200 Principal component analysis (PCA) was performed to verify the intrinsic variation in the data
201 sets. Factor analysis (FA) (Burt, 1950) and general discriminant analysis (GDA) (McLachlan,
202 1992) were used to classify the vinegars according to their NMR fingerprint. To perform
203 GDA, a reduction in variables with respect to complete data sets was necessary. For the
204 complete spectral region data set the number of variables was reduced considering only the
205 signals which presented a factorial weight during FA $> |0.8|$, and the resulting data set with
206 1453 variables and 76 samples was obtained. For the other two data sets, a less severe
207 variables reduction was applied, by simply reducing in spectral resolution. The number of
208 variables in fact was halved, compared to the original data sets, thus ensuring that each peak
209 maintains its shape. After the construction of the models, to evaluate the classification
210 performance, the leave-one out method was used as a validation procedure (Henrion &
211 Henrion, 1994). The most significant signals resulted from GDA were integrated, using the
212 software Amix 3.7.10 (Bruker Biospin GMBH, Rheinstetten, Germany), and used for
213 classification trees analysis (CTA). The aim was to build a discriminant model able to fit
214 adequately the aging process and the type of sample, in order to find a separation between the
215 extra old TBVM and the old TBVM and identify the signals that allow such separation.
216 CTA is used to predict membership of cases or objects in the classes of a categorical
217 dependent variable from their measurements on one or more predictor variables, and, in the
218 past years, it has been successfully used in different areas of healthcare (Harper & Shahani,
219 2002; Ridley et al., 1998; Harper & Winslett, 2004), and food science (Bertelli, Plessi,
220 Sabatini, Lolli & Grillenzoni, 2007; Cirlini, Caligiani, Palla & Palla, 2010). Three different
221 building tree methods were applied to datasets: (a) Discriminantbased Univariate Splits

222 (DUS); (b) Discriminant-based Linear Combination Splits (DLCS); (c) Classification &
223 Regression Tree-style Exhaustive Search for Univariate splits (C&RT). For all of these three
224 building methods, the FACT-style direct stopping was used as stopping rule (Loh &
225 Vanichestakul, 1988). To estimate the prediction capacity of the models, a one-third cross-
226 validation method was applied.

227 All calculations were performed using the PLS_Toolbox version 5.2.2 for Matlab, Statistica
228 6.1 (StatSoft® Italia, Vigonza, Italy) and SPSS 13.0 (SPSS Inc., Chicago, IL, USA).

229

230 **3. Results and discussion**

231 One of the main differences that BVM and TBVM show at macroscopic level is the sugar
232 concentration, which directly influences the density (°Brix). As well-known (Masino,
233 Chinnici, Franchini, Ulrici & Antonelli, 2005), this parameter shows an increase with aging,
234 and this is confirmed also by the density values measured in our samples, reported in **Table 1**.
235 Another important parameter, which correlates density with total acidity, is *R ratio* (Gullo et
236 al., 2004; Satrioni, 2010). For traditional balsamic vinegars produced according to set rules,
237 the optimal value of *R ratio* is between 7–10 (Gullo et al., 2004). The 76 samples show *R*
238 values close to this range, between 8.75 and 12.91 (**Table 1**). Although our results are slightly
239 higher, all the TBVM samples were considered valid and used in the following analyses, since
240 similar *R* values were obtained also by other authors (Consonni, Cagliani, Rinaldini & Incerti,
241 2008).

242 A ¹H NMR spectrum recorded at 400 MHz of a TBVM sample, with expansions of
243 aliphatic/alcoholic, and sugar regions, is shown in **Figure 1**.

244 The sequence allows to suppress broad signals, in our case the water signal, which may be
245 selectively removed, and to enhance narrow resonances. Unfortunately also the large signals
246 of the hydroxyl groups are suppressed; anyway the spectra are very rich in information. This

247 sequence thus provides a sensitive means of investigating the composition of the balsamic
248 vinegar samples object of this study. The metabolites were assigned on the basis of additional
249 2D NMR experiments of samples, by recording NMR spectra of pure compounds, and
250 confirmed by comparing our results with literature data (Cirlini et al., 2009; Consonni,
251 Cagliani, Rinaldini & Incerti, 2008; Consonni et al., 2008; Consonni et al., 2007). The
252 principal metabolites assignments are summarized in **Table 2**. The simple comparison of the
253 calibrated and normalized spectra (**Figure 2**) confirmed that BVM , TBVM old and TBVM
254 extra old show different spectral characteristics, certainly due to the different production
255 techniques, therefore to the consequent different composition. In the case of TBVMs, also the
256 aging process may probably influence the spectral aspect of these samples. The main
257 differences in the spectral intensities distribution are related to the signals of acetic acid,
258 ethanol and acetoin. In TBVM, these compounds seem to be less concentrated in the majority
259 of more aged samples as reported also by other authors. Besides, 5-HMF, fructose and
260 glucose, 6-acetyl glucose and 2,3-butanediol were subjected to larger increase. These
261 compounds have been described in the literature as the most significant ones for monitoring
262 the aging process in TBVs (Caligiani et al., 2007; Masino et al., 2005; Theobald et al., 1998;
263 Cirlini et al., 2009; Consonni et al., 2008). When a comparison between balsamic and
264 traditional balsamic vinegars is performed, it is necessary to specify that decreases and
265 increases in metabolite concentrations are not closely related to the aging process, therefore, it
266 is more cautious, and more correct, to affirm that these differences are probably more related
267 to the intrinsic diversity of products.

268 The PCA was performed on the ^1H NMR complete spectral region to check possible sample
269 grouping (**Figure 3**). This model resulted in 8 PCs explaining 96.91% of the total variance,
270 and it was able to discriminate only the BVMs from TBVMs, without any differentiation
271 between old and extra old TBVMs. This result demonstrated the considerable complexity of

272 the system; therefore, in order to give a clearer interpretation of PCA model, and to verify
273 whether the application of a supervised multivariate statistical analysis was able to classify
274 the samples, GDA was applied. The analysis was performed on complete and also on
275 expanded spectral region data sets, since a less severe variables reduction was necessary in the
276 aliphatic/alcoholic and sugar regions. The best results were obtained using the forward
277 stepwise procedure. The GDA model obtained from the complete spectra was able to group
278 the samples in three evident clusters, corresponding to the type of sample (**Figure 4**). The first
279 two canonical discriminant function (DF) explains 98.6% of the total variance, and the results
280 of the leave-one out cross-validation show a predictive capacity of 98.4%. The first two DFs
281 are particularly correlated with the signals of 5-HMF (C1H, C3H, C4H), α -glucopyranose
282 (α C1H), malic acid (C3H), a signal at 3.3 ppm referred to the glucose and fructose region,
283 succinic and acetic acids. These results were confirmed by the models obtained by analyzing
284 the expanded spectral regions, and were both able to group the vinegars in three evident
285 clusters (score plots not shown). The first DFs of aliphatic/alcoholic and sugar regions data
286 sets explain 98.3% and 89.7% of the total variance respectively, and the results of the leave-
287 one out cross-validation show a predictive capacity of 90.5% and 96.8% respectively. The
288 first two DFs of each model are particularly correlated with the signals of succinic acid, malic
289 acid (C2H), 6-acetyl glucose and the unknown signal at 1.74 ppm for the aliphatic/alcoholic
290 region data set, α -glucopyranose (α C1H), 5-HMF (C6H₂), tartaric acid, β -glucopyranose
291 (β C1H), β -fructopyranose (C6H), 2 signals at 3.3 ppm and 3.03 ppm referred to the glucose
292 and fructose region for the sugar region data set. The signals, corresponding to the most
293 significant compounds in GDA, were integrated and used to build a new CTA discriminant
294 model, in order to evaluate their effects on the balsamic vinegars discrimination. The obtained
295 results are summarized in **Table 3**. In general, when the results of CTA were judged, the
296 number of terminal nodes must be considered and a tree with a small number of terminal

297 nodes must be preferred if the same capacity of classification is reached by different
298 approaches. Considering the classification results and the number of terminal nodes, C&RT
299 seems to be a good compromise between classification capacity and the complexity of the tree.
300 In **Figure 5** the best tree obtained by C&RT method is reported. As evident, the model is able
301 to discriminate between the extra old TBVM and the old TBVM and to identify the signals
302 that allow such separation. The most interesting finding resulted from CTA is that in this
303 model the discrimination between old TBVM and extra old TBVM is essentially due to the
304 signals of acetic acid, 5-HMF (C4H), malic acid (C3H), β -glucopyranose (β C1H),
305 corresponding to those compounds already identify as significant ones also in the literature
306 (Caligiani et al., 2007; Consonni et al., 2004; Theobald et al., 1998; Consonni et al., 2008),
307 and to the unknown compound at 1.74 ppm.

308

309 **4. Conclusion**

310 The NMR spectroscopy, coupled with multivariate analysis, has demonstrated to be a
311 powerful tool in BVM and TBVM characterization and quality control. Statistical analysis
312 showed that the signals of 5-HMF, α and β -glucopyranose, malic, succinic, tartaric and acetic
313 acids, 6-acetyl glucose, a signal at 3.3 ppm referred to the glucose and fructose region were
314 the most statistically significant variables. All these compounds are described in the literature
315 as relevant ones for discriminating the balsamic vinegars and for monitoring the aging process.
316 Theobald et al. (1998) established that the highest concentration of 5-HMF was found in TBV
317 samples. The changes in sugars and in acetic acid contents occurring during the aging process
318 of TBVs are a well known finding (Caligiani et al., 2007; Consonni et al., 2008). In 2004, as a
319 part of NMR studies on Italian balsamic and traditional balsamic vinegars, Consonni et al.
320 (2004) confirmed that, among several compounds, the highest content of malic acid was
321 found for the older BV and TBV, while much lower values were found for the younger BVs.

322 Cocchi et al. (2002) proved that succinic acid increases in the young vinegars and decreases in
323 the old ones. The hypothesis of formation of glucose and fructose acetates during maturation
324 and aging of balsamic vinegar was verified by Cirlini et al. in 2009.

325 All these findings provide interesting additional knowledge on TBVs, usable to reinforce and
326 safeguard the wealth of Modenese traditions, represented by these unique products.

327

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335

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471 **Figure Captions**

472 **Figure 1.** Typical ^1H NMR spectra recorded at 400 MHz of a TBVM sample; expansions of
473 (A) aliphatic/alcoholic, and (B) sugar regions with metabolite assignments (see Table 2).

474 **Figure 2.** Comparison of the ^1H NMR spectra calibrated and normalized using the TMS
475 signal of BVM (a), TBVM old (b) and TBVM extra old (c).

476 **Figure 3.** PCA score plot performed on complete spectral region data set: TBVM extra old
477 (Δ), TBVM old (+), and BVM (\square).

478 **Figure 4.** Vinegar classification using GDA on complete spectral region data set, showing the
479 separation of samples in three evident clusters: TBVM extra old (Δ), TBVM old (+), BVM (\square)
480 and group centroids (\bullet).

481 **Figure 5.** Vinegar classification using CTA (C&RT-style method) on the data set obtained by
482 integrating the most significant signals resulted by GDA.

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Table 1. Values of °Brix and *R ratio* in TBVM and °Brix in BVM samples

TBVM samples	Type^a	°Brix	<i>R ratio</i>^b	BVM samples	°Brix
1	Old	65	10.50	41	32
2	Old	67	12.91	42	22
3	Old	61.5	9.70	43	15.25
4	Old	69.5	11.94	44	26.5
5	Old	68.25	9.50	45	24
6	Extra old	71	10.88	46	19
7	Extra old	71.75	11	47	27
8	Extra old	71	11.14	48	23.5
9	Extra old	73	11.09	49	15.3
10	Extra old	70.5	12.56	50	20
11	Old	68.5	11.01	51	19.5
12	Extra old	73	9.92	52	19
13	Extra old	74	12.07	53	29.5
14	Old	70	11.59	54	38.5
15	Extra old	72	9.28	55	32
16	Extra old	71	11.75	56	36.5
17	Old	60.5	8.75	57	44.5
18	Old	65.5	9.86	58	40
19	Old	68	10.33	59	28.5
20	Extra old	72	12.16	60	39
21	Extra old	72.5	10.63	61	38.5
22	Extra old	70	10.08	62	53.5
23	Extra old	73	10.08	63	38.5
24	Extra old	72.5	10.54	64	42.5
25	Extra old	71.75	11.21	65	36
26	Old	65	9.88	66	18
27	Old	63	9.36	67	36
28	Old	64	10.24	68	31
29	Old	63.5	9.92	69	29
30	Old	65	10.25	70	30
31	Extra old	71.5	10.87	71	38.5
32	Extra old	71	9.89	72	53
33	Extra old	73.5	11.38	73	19
34	Extra old	72	11.41	74	38
35	Extra old	71	10.64	75	25
36	Extra old	72.5	11.49	76	30
37	Extra old	70.5	10.72		
38	Extra old	71	8.75		
39	Old	60.75	12.90		
40	Old	62	9.92		

^aAge is indicated only for known aging process samples. Extra old >25 years; old >12 and <25 years. For BVM the aging is unknown, however it is < 3 years.

^b°Brix/Total acidity % (p/v).

Table 2. Metabolites and ^1H chemical shifts identified^a

Compound	Group	δ (ppm)	Multiplicity ^b	J (Hz)
Acetic acid	C2H ₃	1.90	s	-
Acetoin	C4H ₃	1.13	d	7.0
	C1H ₃	2.08	s	-
2,3-Butanediol	C1H ₃	0.93	d	5.9
	C4H ₃	0.98	d	5.9
Ethanol	C2H ₃	1.04	t	7.2
Formic acid	HCOOH	8.16	s	-
β -Fructofuranose	C1H	3.44	m	-
	C6H	3.40	m	-
β -Fructopyranose	C1H	3.32; 3.48	dd	11.2; 5.4
	C3H C4H	3.62	m	-
	C5H	3.68	m	-
	C6H	3.82	d	12.0
6-Acetyl glucose signals ^c	CH ₃ CO	1.93-2.01	m	-
α -Glucopyranose	α C1H	4.89	d	3.5
	C4H	3.09	m	-
	C6H	3.62	m	-
β -Glucopyranose	β C1H	4.26	d	7.9
	C2H	2.87	t	8.4
5-HMF	C1H	9.56	s	-
	C3H	7.52	d	3.5
	C4H	6.63	d	3.3
	C6H ₂	4.47	s	-
Lactic acid	C3H ₃	1.19	d	6.7
Malic acid	C2H	2.38	m	-
	C2'H	2.59	m	-
	C3H	4.07	m	-
Succinic acid	C2H ₂ C3H ₂	2.40	s	-
Tartaric acid	C2H C3H	4.31	s	-
Valine	C γ H ₃	0.87	d	6.8

^aAssignments were from HSQC and HMBC experiments. The chemical shifts were expressed as relative values to those of TMS at 0 ppm.

^bPeak multiplicities: s, singlet; d, doublet; t, triplet; dd, doublet of doublets; m, multiplet.

^cEsters of glucose (6-acetylglucose) in the two anomeric forms. See Cirlini *et al.* for more details.³²

Table 3. Calibration and validation results using CTA with building tree methods on the data set obtained by integrating the most significant signals resulted by GDA

Building tree method	All samples calibration (n=76)		One-third validation (n=50+26)					
	<i>Terminal nodes</i>	<i>Correct classification (%)</i>	Calibration set		Validation set			
			<i>Terminal nodes</i>	<i>Correct classification (%)</i>	<i>Correct classification (%)</i>			Total
				1	2	3		
DUS	23	100	12	100	75	62	100	81
DLCS	5	100	4	100	0	62	100	67
C&RT	12	100	9	100	100	50	100	81

Figure 1

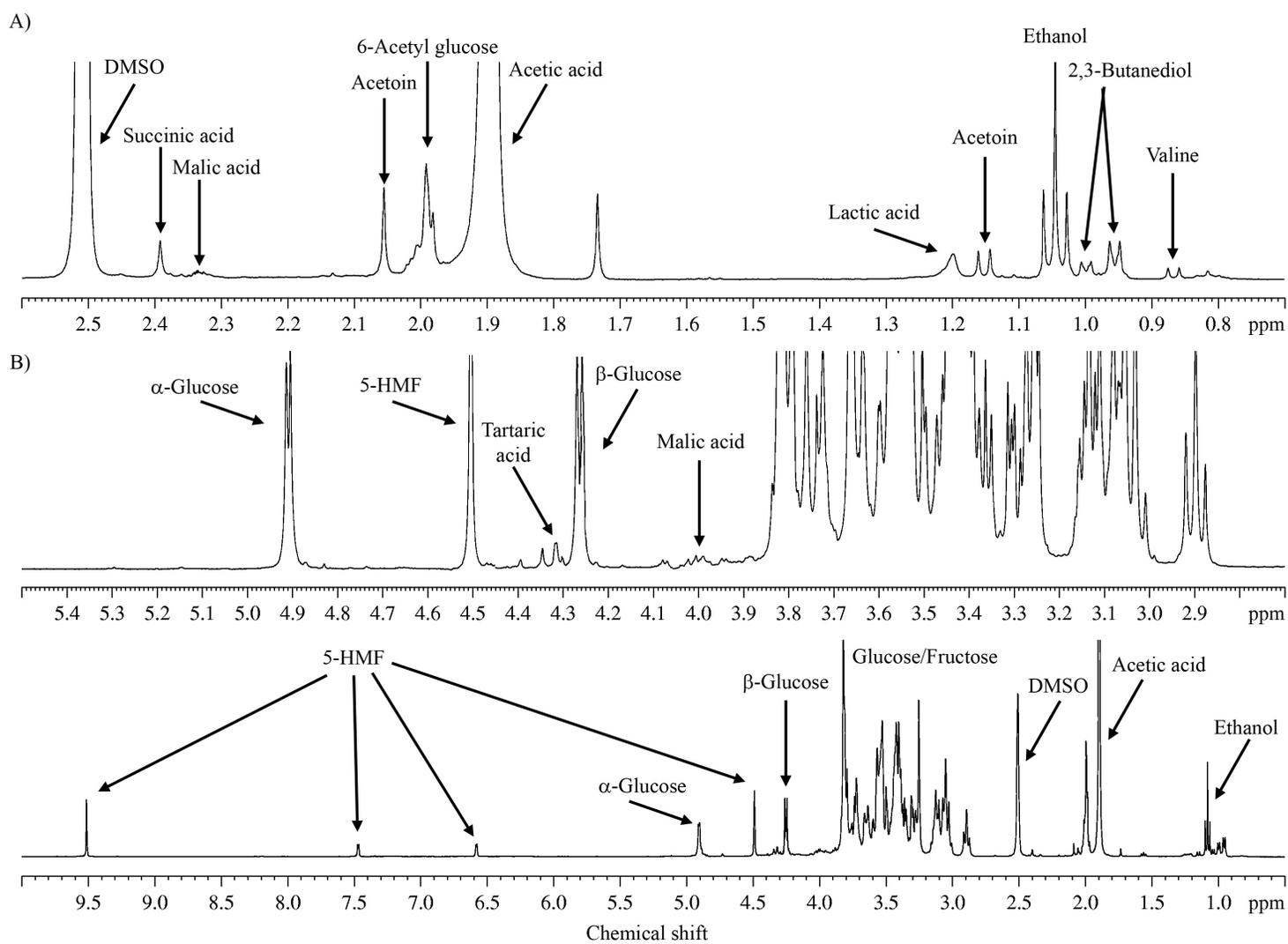


Figure 2 color for web

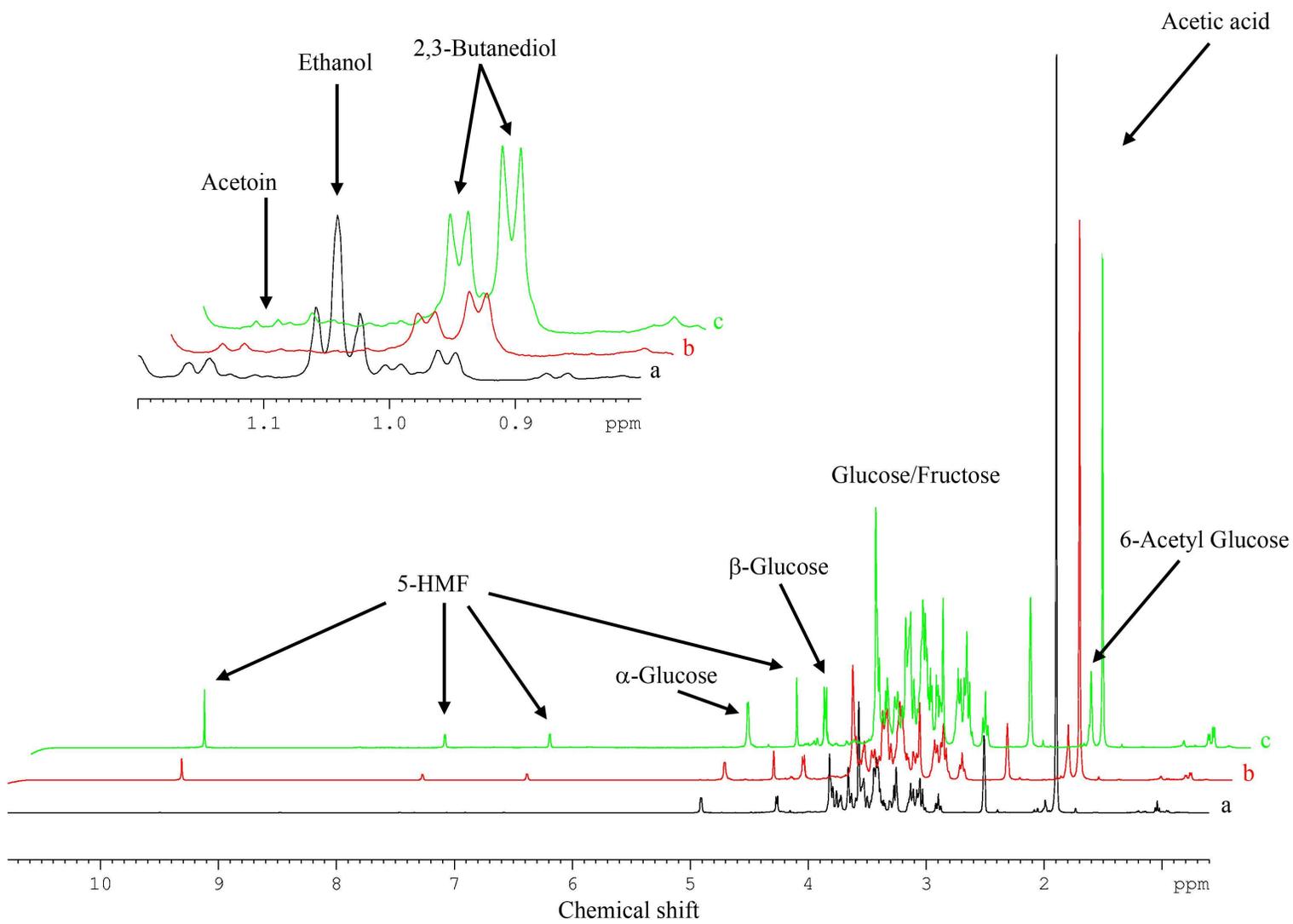


Figure 2 bw for print version

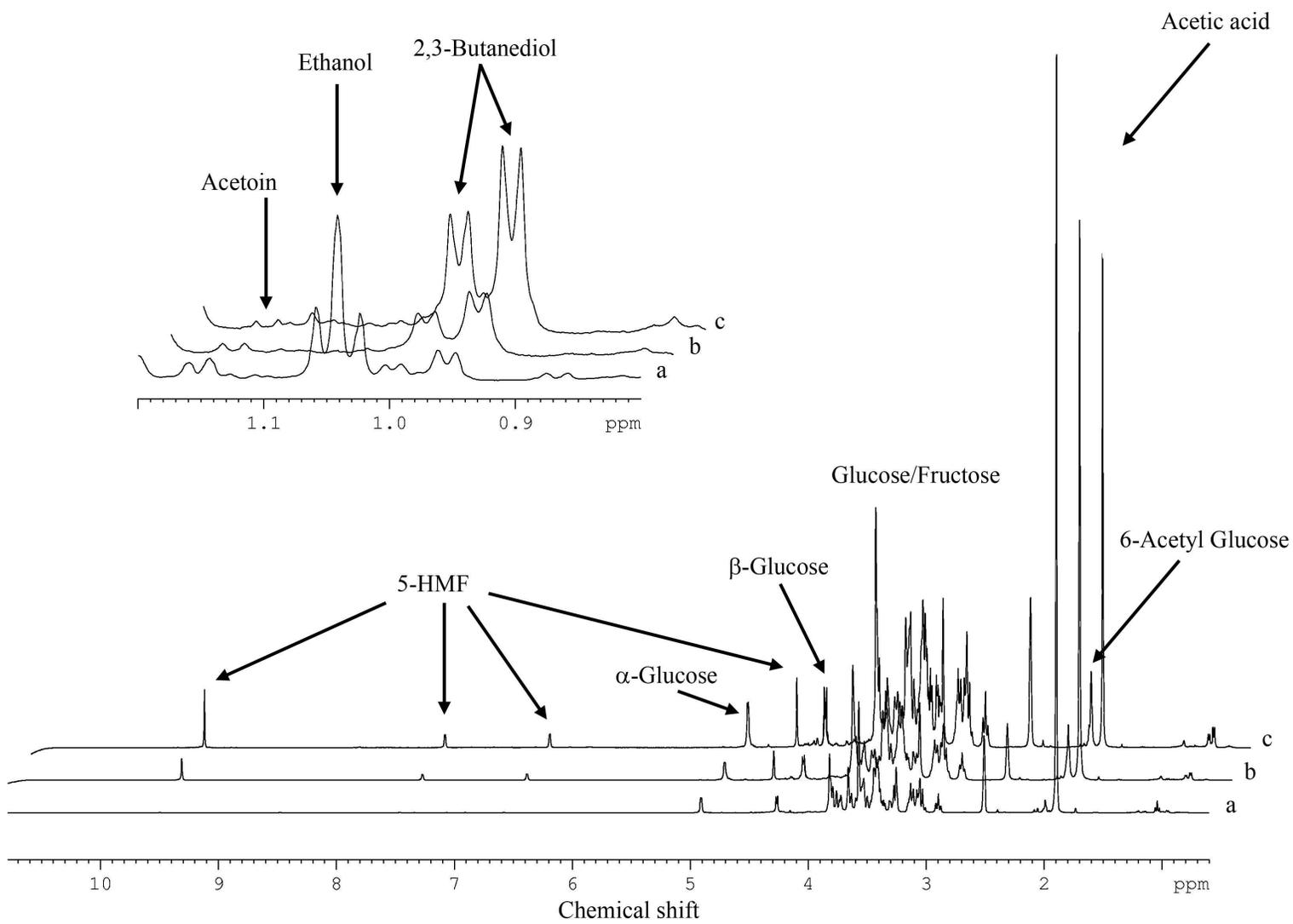


Figure 4

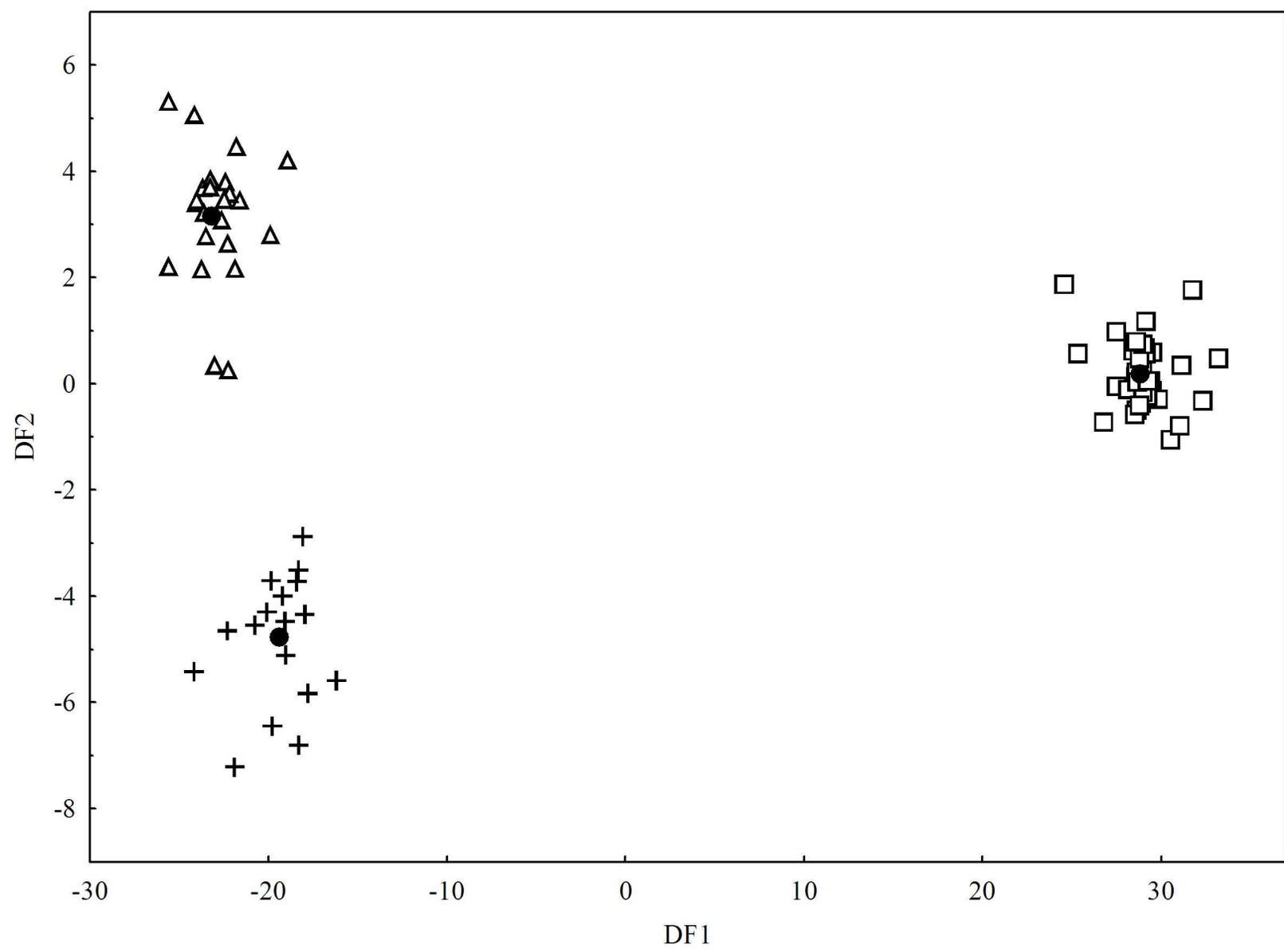


Figure 5 color for web

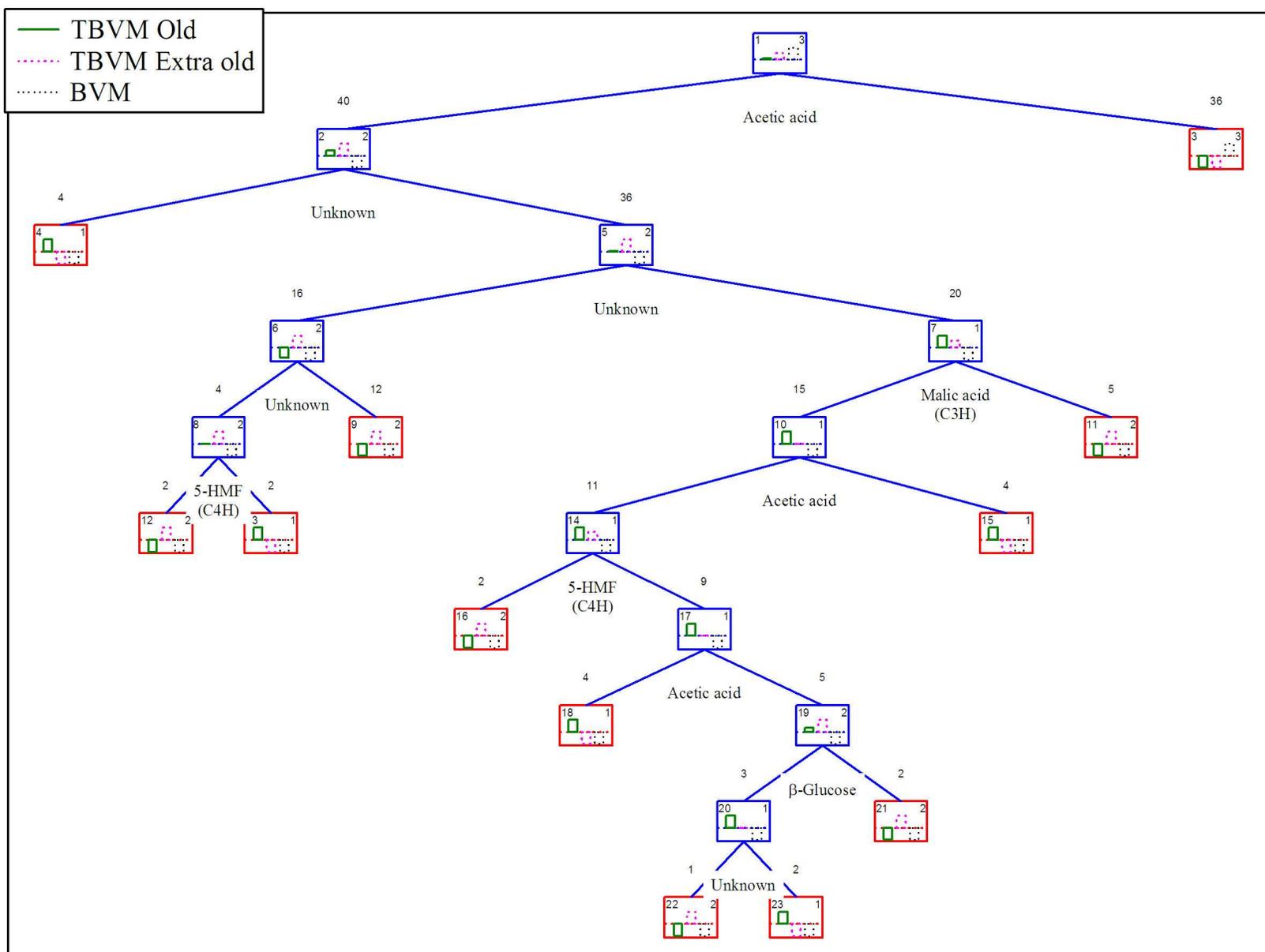
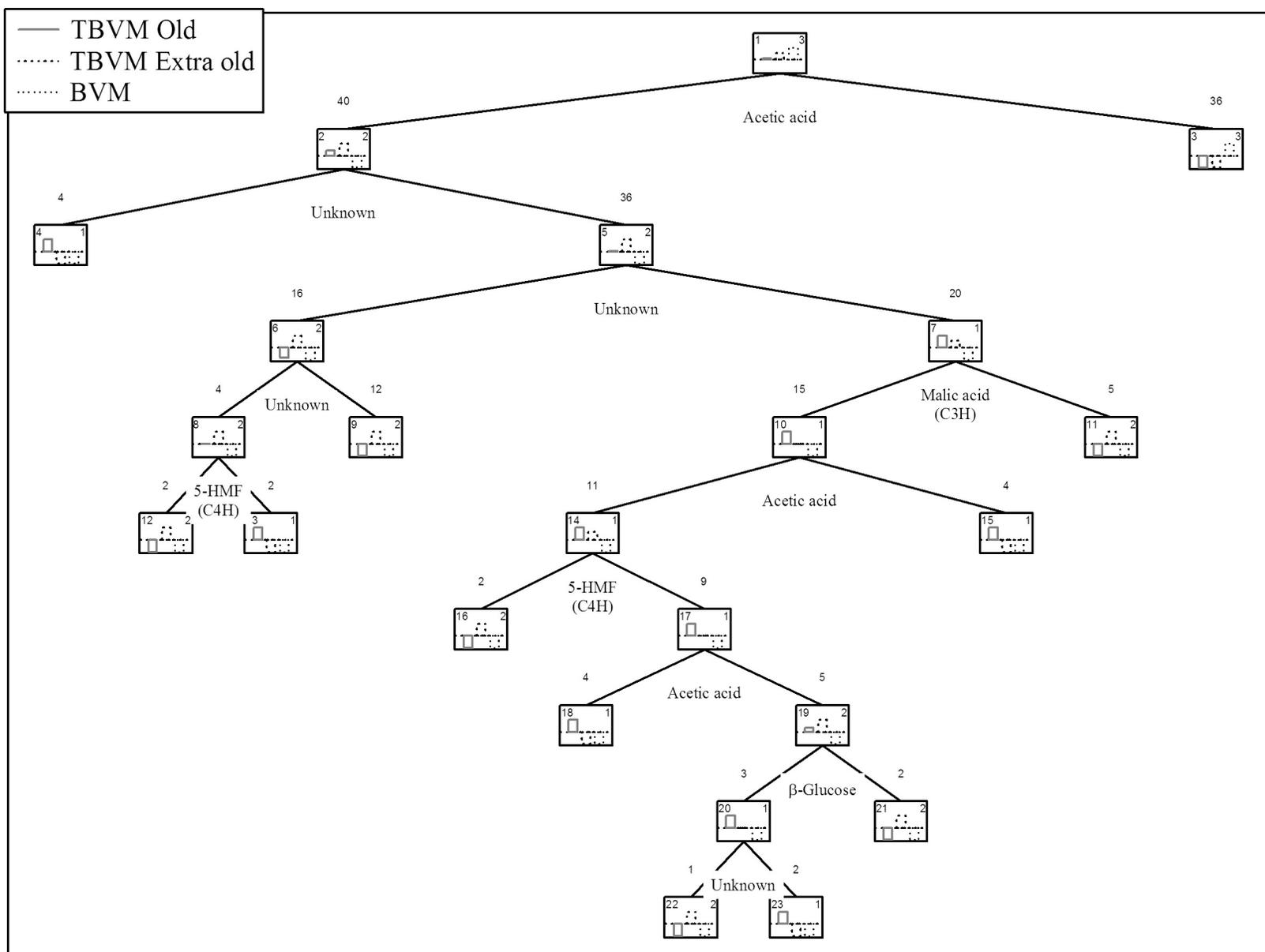


Figure 5 bw for print version



LWT-Food Science & technology

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Traditional Balsamic vinegar and Balsamic vinegar of Modena analyzed by Nuclear Magnetic Resonance spectroscopy coupled with multivariate data analysis

Dear editor,

The manuscript was revised according to the comments

1. Number of keywords provided should be within 4-5.
 - Keywords are now 5;
2. Research papers should be within 20 pages of double-spaced type, (not including figures/tables and their captions).
 - The manuscript was revised and shortened to 19 pages;
3. Avoid vertical lines in tables.
 - The tables were corrected and vertical lines were deleted;