

1 Research Paper

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3 **Non-targeted Spatially Offset Raman Spectroscopy-based vanguard analytical**  
4 **method to authenticate spirits: White Tequilas as a case study**

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18 *Abstract*

19 Adulteration and counterfeiting are ongoing problems for alcoholic drinks, including beers,  
20 wines, and spirits. To fight against them, official analytical methods need to be complemented  
21 with faster, trustworthy, non-invasive and *in-situ* ones, which have been named as vanguard  
22 methods, to increase the efficiency in the detection probability of truly adulterated alcoholic  
23 drinks. The analytical methodology proposed here synergistically combines a novel  
24 measurement analytical technique (spatially offset Raman spectroscopy, SORS) with  
25 chemometrics methods, i.e., principal component analysis (PCA), soft independent modeling  
26 of class analogies (SIMCA), partial least squares regression-discriminant analysis (PLS-DA),  
27 support vectors machine, (SVM) and quantitative partial least squares regression (PLSR).  
28 The applicability of the proposal is tested with Tequila to (i) differentiate among *100% agave*  
29 and *mixed* white packaged Tequilas, and (ii) to predict the alcoholic content. SORS spectra of  
30 51 samples were obtained in the 300-2000  $\text{cm}^{-1}$  range, from which classification and  
31 regression models were developed. The best classification performances were obtained with  
32 PLS-DA and SVM with 100% sensitivity, specificity and overall classification rate. PLSR  
33 exposed a better trend of the samples than PCA in the exploratory analysis; and yielded  
34 predictive models capable of foreseeing alcoholic contents with average errors lower than 4%.  
35 These results demonstrate the potential of this fast, *in-situ* analytical approach to be used as a  
36 vanguard analytical method to screen adulterated or counterfeited Tequilas and to assess the  
37 conformity of the alcoholic stated in the label.

38

39 *Keywords*

40 Chemometrics; Spirits fraud; Spatially offset Raman spectroscopy; Tequila authentication.

41

## 42 ***1. Introduction***

43 Criminal activity against consumers continues non-stop, in fact European Union Intellectual  
44 Property Office (EUIPO) and European Union Agency for Law Enforcement Cooperation  
45 (EUROPOL) have indicated in a last report published in March 2022 that *the production of*  
46 *illicit food products, especially drinks, is increasingly professional and sophisticated* [1].

47 However, in terms of health and food safety, the weightiness of food and drink fraud will  
48 depend on the type of fraud. In some cases, the consequences are limited to consumer  
49 deception, since offenders pass off lower value products as higher value foods or drinks for  
50 illicit financial profit. Specifically in drinks the most frequent fraud is that committed in  
51 alcoholic beverages, so-called spirits. In fact, in the last two years, adulteration of this type of  
52 product has been detected, such as the case of the Whiskey fraud in Spain in 2020 [2] or the  
53 adulteration of alcoholic beverages in Santo Domingo in April 2022, which resulted in the  
54 death of several people [3].

55 There is a battery of recognized and well-described analytical methods for detecting different  
56 types of adulteration for each particular alcoholic beverage, most of them based on the  
57 identification and quantification of specific chemical markers. Despite traditional analytical  
58 methods proved to be reliable, accurate and are suitable tools for production control, they  
59 often do not comply with the principles of green chemistry, since they involve the use of  
60 environmentally unfriendly reagents, are time-consuming and frequently expensive,  
61 considering them as rearguard methods [4]. This gives opportunity for the development and  
62 application of alternative analytical methods, which are characterized by being miniaturized,  
63 transportable, simple, rapid, low-cost and capable of providing overall analytical information  
64 that is reliable and representative. The application of these type of alternative analytical  
65 methods, which have been named as vanguard methods, increase the efficiency of control  
66 laboratories since they make possible the analysis of only suspicious samples by rearguard

67 methods [4]. The term vanguard method does not refer to the fact that the methodology  
68 presented in this study is highly recent and innovative, as might be inferred at first. It suggests  
69 that such a methodology could be applied as a first analytical approach to quickly process  
70 laboratory samples. In this sense, a vanguard method is often a forward screening method that  
71 allows the selection of suspect samples that will subsequently be subjected to a full backward  
72 analytical method, *i.e.*, a reanguard method.

73 In this sense, the use of non-targeted spectroscopic analytical techniques, such as  
74 conventional Raman or medium and near infrared spectroscopies, constitute established  
75 methodologies that fit most requirements to get vanguard analytical methods as they require  
76 minimum or null sample preparation. Despite of providing unspecific signals (spectroscopic  
77 instrumental fingerprints), they became popular to determine the composition/adulteration of  
78 food and beverages to ensure the authenticity and traceability [5]. One essential and inherent  
79 subsequent step after the application of spectroscopic techniques is the use of multivariate  
80 chemical data analysis or chemometrics, which together have created a synergistic and  
81 powerful analytical methodology that is regularly applied in the food industry to extract  
82 important and non-evident (or hidden) information from the raw spectra by developing  
83 mathematical models [6,7,8].

84 Quite recently, a new and more advanced Raman spectroscopy modality, termed spatially  
85 offset Raman spectroscopy (SORS), appeared and it shows highly promising capabilities for  
86 spirit quality and authenticity control. The fundamentals of SORS are like the conventional  
87 Raman spectroscopy, although in SORS the Raman signal is obtained at certain millimeters  
88 off the laser spot, making it possible to collect photons emitted from samples contained within  
89 opaque packaging materials [9]. This means that it is possible to carry out the analysis directly  
90 on the product within the container, without the need to alter the original package/sample,  
91 making SORS one of the few truly non-invasive analytical techniques. Even though this novel

92 approach was first developed for the pharmaceutical industry, it expanded rapidly to the food  
93 industry to analyze packaged beverages in a fast and non-destructive manner [9]; for instance:  
94 Vodka, Gin and Whisky through their containers [10]. However, no applications have been  
95 found to authenticate Tequilas.

96 Tequila is a representative spirit from México that holds an Official Designation of Origin  
97 (DOT - *from the Spanish term 'Denominación de Origen Tequila'*), which is regulated by the  
98 Mexican Government and the Regulatory Council of Tequila (CRT) through the official  
99 Mexican standard NOM-006-SCFI-2012 [11]. Tequila can be classified in five classes  
100 according to their aging process in oak or holm oak containers: '*Silver or White*', '*Aged*',  
101 '*Extra-aged*' and '*Ultra-aged*' according to whether maturation lasts for <2 months,  $\geq 2$   
102 months,  $\geq 2$  years or  $\geq 3$  years, respectively. '*Gold Tequila*' corresponds to commercial  
103 mixtures of White Tequila with Aged, Extra-aged or Ultra-aged Tequilas [11]. Additionally,  
104 two categories of Tequila can be distinguished: (i) 100% agave Tequila if only sugars from  
105 the juice of the *Agave Tequilana Weber blue variety* are used for the fermentation process,  
106 and (ii) 'mixed Tequilas' if any combination with other sources of reducing sugars (never  
107 more than 49%) are added to the process. The commonest commercial product is white  
108 Tequila, so, this paper focused on it.

109 Currently, many adulteration and counterfeiting cases are still reported, not only at Mexico  
110 but in other countries. The main adulteration practice is to substitute ethanol with methanol  
111 or, less frequently, with propanol, ethylene glycol, aldehydes and others [12]. In 2021, a  
112 production of 527 million of liters of Tequila was reported by the CRT whose quality and  
113 authenticity were evaluated using representative samples extracted from the distilleries and  
114 analyzed independently at the CRT. All the aforementioned classes of Tequila are inspected  
115 by the CRT using standardized analytical techniques, such as liquid and gas chromatography  
116 or atomic absorption spectroscopy, to adhere to current official analytical methods. Several

117 quality parameters are determined, e.g., furfural, esters, aldehydes, methanol, higher alcohols,  
118 reducing and total sugars. An exemplary routine verification is whether the alcoholic content,  
119 using a digital densimeter method at 20°C, which is established in the Mexican standard  
120 NMX-V-013-NORMEX-2019 [13], is between 35 and 55% (v/v).

121 The studies found in the literature concerning the assessment of tequila authenticity are  
122 focused on (i) some chemical markers, (ii) a specific spectral region of interest (ROI), or (iii)  
123 Red, Green and Blue (RGB) color coordinates obtained after the Tequila analysis by  
124 chromatographic and spectroscopic analytical techniques [14,15,16,17,18,19]. For example,  
125 Contreras et al. [20] applied UV-Vis spectroscopy to identify adulterated and fake Tequilas  
126 (between white and rested tequila) or Perez-Beltran et al. [21] employed FTIR and data fusion  
127 approach for distinguishing between pure and mixed White Tequilas. However, surprisingly  
128 no studies have been found where the full RAMAN spectrum is used as an unspecific  
129 instrumental fingerprint but characteristic of each tequila together with chemometric tools for  
130 tequila authentication.

131 In this regard, the innovation of this work lies in developing a fast and non-invasive vanguard  
132 analytical method for the *in-situ* screening quality control of spirits using SORS. Its  
133 applicability is demonstrated to ensure Tequila from Mexico in the following terms: (i)  
134 discriminate White Tequilas (100% agave vs mixed), and to (ii) predict and verify the  
135 alcoholic content. For this, SORS spectra were used together chemometric tools to develop  
136 suitable classification and quantitation multivariate analytical methods. Classification  
137 methods were validated in terms of sensitivity, specificity, precision, negative prediction  
138 value, among other 21 classification performance metrics and estimated following the study  
139 published by Cuadros-Rodríguez et al. (2016) [22]. In addition, the quantitative method for  
140 determining the alcohol content was validated according to the ASTM E2617 standard [23].

141

142 **2. *Materials and methods***

143

144 **2.1. *Tequila samples***

145 A total of 51 White Tequila samples were provided by the CRT in México, and analyzed in  
146 Spain, as described in the 'spatially offset Raman spectroscopy (SORS) measurements'  
147 section. Thirty White Tequilas belonged to the 100% agave White Tequila category (TB -  
148 from the Spanish term 'Tequila Blanco') and twenty-one to the mixed White Tequilas (TBM -  
149 from the Spanish term 'Tequila Blanco Mixto'). The alcoholic content of all these samples was  
150 determined by the CRT using a digital densimeter at 20°C [13].

151

152 **2.2. *Spatially offset Raman spectroscopy (SORS) measurements***

153 Vaya Raman SORS equipment (Agilent Technologies, Santa Clara, CA, USA) was used. The  
154 excitation radiation was 830 nm with a maximum power laser of 450 mW, obtaining Raman  
155 spectra in the low frequencies range, from 350 to 2000  $\text{cm}^{-1}$ , with 12 to 20  $\text{cm}^{-1}$  spectroscopic  
156 resolution. The SORS measurements of the 51 white Tequila samples were performed directly  
157 through amber vials lasting 30 s, approximately.

158

159 **2.3. *Similarity analyses***

160 In order to make sure that this methodology can be transferable to any other situation,  
161 similarity analyses were performed. SORS measurements were directly performed on four  
162 original bottled Tequilas marketed in Spain (2 mixed White Tequilas, 1 mixed Rested Tequila  
163 and 1 mixed Tamarind flavored White Tequila). Afterwards, 2 mL of each of them were  
164 transferred to amber glass vials, similar to those used to transport the Mexican Tequila  
165 samples, and measured. Once both spectra for each sample were acquired, the similarity  
166 among them was assessed calculating the corresponding nearness similarity index [24], which

167 is based on the proximity of two vectors in space and is calculated from the standardized  
168 Euclidean distance, as depicted in Eq. (1).

$$169 \quad \text{NEAR}(X_{\text{SORS}}, X_{\text{CRS}}) = 1 - \left[ \frac{(X_{\text{SORS}} - X_{\text{CRS}}) \times (X_{\text{SORS}} - X_{\text{CRS}})^{\text{T}}}{(X_{\text{SORS}} + X_{\text{CRS}}) \times (X_{\text{SORS}} + X_{\text{CRS}})^{\text{T}}} \right] \quad (1)$$

170 where  $X_{\text{SORS}}$  and  $X_{\text{CRS}}$  symbolized both SORS and conventional Raman spectra, respectively,  
171 and the superscript T denotes the transposed matrix [25].

172

#### 173 ***2.4. Multivariate data analyses***

174 SORS raw data were exported from CSV format (comma-separated values) to MATLAB  
175 environment (Mathworks, Massachusetts, USA, v. R2013b). The exported spectra contained  
176 1651 variables, each. The training set was constituted by 41 samples (24 of TB type and 17 of  
177 TBM type) whilst the external validation set contained 10 different samples (6 TB and 4  
178 TBM). Splitting was performed applying the Kennard-Stone selection method (so-called  
179 CADEX algorithm), which was deployed on the TB and TBM classes independently in order  
180 to select the samples of the validation set.

181 The multivariate data analyses were carried out using the PLS\_Toolbox software (v. 8.6.1,  
182 2019, Eigenvector Research In., Manson, WA, USA). The applied chemometric tools were  
183 principal component analysis (PCA) and partial least squares regression (PLSR) for  
184 exploratory analysis, soft independent modeling of class analogy (SIMCA), partial least  
185 squares-discriminant analysis (PLS-DA) and support vector machines (SVM) for  
186 classification, and PLSR was also used to quantify the alcoholic content of the samples. Mean  
187 centering and smoothing were used as pre-processing techniques depending on the  
188 multivariate method, as described in 'exploratory analyses' and 'classification analyses'. The  
189 proper number selection of the PCs and LVs of the models was based on the study of their



190 root mean square error for calibration (RMSEC), or for prediction (RMSEP) and for cross-  
191 validation (RMSECV) plots, and the total explained variance, avoiding overfitting in each  
192 case.

193

### 194 **3. Results and discussion**

195

#### 196 **3.1. SORS analyses and characterization**

197 When SORS analyses are performed, two measurements are acquired: one at zero offset and  
198 another one with a laterally spatial offset of 0.7 mm from the point of incidence of the laser to  
199 the collection point [9]. This separation favors the photons from the lower layers to be  
200 radiated from a spot laterally shifted from the incidence zone while the photons on the upper  
201 package are radiated from the same incidence zone [26]. Afterwards, internal pre-processing  
202 and normalization are performed by the equipment and a final Raman spectrum is obtained  
203 with no contribution of the container. The Raman spectra of the two categories of white  
204 Tequilas can be observed in Fig. 1.

205

*Fig. 1*

206

207 The intense peak located at  $882\text{ cm}^{-1}$  and the peak at  $1053\text{ cm}^{-1}$  are attributed to the stretching  
208 and deformation modes of the skeletal C-C-O moieties, whilst the peak at  $1090\text{ cm}^{-1}$  is  
209 associated to the stretching mode of the C-O bond. The peaks at  $1279\text{ cm}^{-1}$  and  $1455\text{ cm}^{-1}$  are  
210 assigned to the deformation wagging mode and to the wagging mode of  $\text{CH}_2$ , respectively  
211 [15,27]. Additionally, the two small peaks around  $1610\text{ cm}^{-1}$  and  $1728\text{ cm}^{-1}$  are associated to  
212 the cyclic ketone structure, which is the basis of furanic compounds in Tequila. Noteworthy,

213 those peaks are more intense for the TB category than for the TBM one, as TB proceeds only  
214 from fermentable sugars of the Agave Tequilana Weber blue variety (through the Maillard  
215 reaction [28] when cooked). On the contrary, TBM might or might not present these spectral  
216 Raman peaks because this category of Tequilas can be produced from mixtures of fermentable  
217 sugars, so that the production of furanic compounds might not occur [29].

218 These acquired signals (Raman spectra), which are here used to evaluate the authenticity and  
219 quality of White Tequilas, are non-specific instrumental fingerprints and make it necessary  
220 the application of multivariate data analyses, as described in the following subsections.

221

### 222 *3.2. SORS and conventional Raman spectra similarity analyses*

223 A point-by-point comparison, using the nearness similarity index (NEAR), among the four  
224 pairs of spectra (data vectors) corresponding to the Tequila samples marketed in Spain was  
225 performed to assess their similarity when the spectroscopic measurements are performed  
226 through the original Tequila glass bottle or through amber glass vials (used as reference). The  
227 expected NEAR results of the standardized Euclidean distance range from 0 to 1, being 1 the  
228 maximum similarity among the spectra. Fig. 2 displays the spectra of the four analyzed  
229 samples within their original glass bottles and the spectra of the samples transferred to the  
230 vial.

231

Fig. 2

232

233 As it can be observed in Fig. 2, each pair of overlapping spectra are similar at first glance and  
234 this fact is further confirmed when the Nearness similarity index is calculated, obtaining  
235 NEAR values  $>0.92$ , which indicates that both spectra are largely similar with almost null

236 influence of the original glass bottles over the measurements (the remaining ca. 0.08% can be  
237 considered as random noise). According to these results, it is evident that the methodology  
238 presented here has potential application to the *in-situ* quality control and authentication  
239 analysis of Tequila.

240

### 241 **3.3. Exploratory analyses**

242 An exploratory analysis was performed to screen the natural grouping of the 51 Tequilas. For  
243 this study, the spectral data was previously mean centered. First, a PCA was built considering  
244 5 principal components (PCs) and explaining 75.9% of the cumulative variance, whose main  
245 scores plot, is displayed in Fig. 3. Nonetheless, it can be observed that the samples do not  
246 follow any specific trend among categories.

247

Fig. 3

248

249 Furthermore, PLSR was used to explore these samples. The model was built with 5 latent  
250 variables (LV) explaining 71.1% of the cumulative variance in the X block and 85.8% in the  
251 Y block. Fig. 4 shows the LV2 vs LV3 scores plot, where the TB category concentrates  
252 (although not unequivocally) in the upper-right region of the plot and the TBM category to  
253 the left. The different results among PCA and PLSR lies basically in the very nature of the  
254 PLS latent variables that capture both variance and correlation [30], yielding best results when  
255 PLSR is applied, as it was also found when looking for groups among FTIR fused data of  
256 100% agave and mixed White Tequilas [21]. Additionally, there are some samples placed out  
257 of the 95% confidence limit that might be considered as outliers (see Figures 3 and 4),  
258 however, it was noticed through the normalized (or reduced) Hotelling  $T^2$ -leverages vs. Q

259 residuals plot that those samples had a normal behavior, discarding the existence of outliers.  
260 Thus, all samples were included in the following data analyses.

261

Fig. 4

262

### 263 **3.4. Classification analyses**

264 The next step after the exploratory analysis was the development of non-targeted multivariate  
265 analytical methods to discriminate among TB and TBM. For all classification models, mean  
266 centering and smoothing (Savitski-Golay, 15 points for filter width and 1<sup>st</sup> order polynomial)  
267 were used as preprocessing techniques. Smoothing is a low-pass filter that removes high-  
268 frequency noise [30]. The target class is TB as it is the category with more probability to be  
269 adulterated due to its economic profit. The results of the final classification models are  
270 discussed next.

#### 271 ■ One Class-SIMCA

272 The developed SIMCA models were generated using two strategies: (i) two input-class  
273 classification (2iC-SIMCA) models, in which the model is trained using two classes (TB  
274 and TBM), and (ii) one input-class classification (1iC-SIMCA) model, in which the  
275 model is trained only with the 'target class' (TB). Within the 1iC-SIMCA strategy, two  
276 options were evaluated: (a) using the aforementioned calibration and validation data sets  
277 and (b) augmenting the validation set using all the 21 TBM and the previous 6 TB  
278 samples. It was found that the 1iC-SIMCA approach presented the best results using 5  
279 PCs.

280 The 1iC-SIMCA classification plot (Fig. 5a) depicts the normalized (or reduced)  
281 Hotelling's  $T^2$  and Q statistics of the target class, at a 95% confidence level. Samples

282 from the validation set with normalized  $T^2$  and  $Q$  values  $< 1$  (left-bottom quadrant) are  
283 those considered as the target class (TB), whereas samples with  $T^2$  and  $Q$  values  $> 1$   
284 (right-bottom quadrant) are considered as non-TB (or TBM). In this sense, samples  
285 TBM13 and TBM102 are misclassified as TB and sample TB70 as TBM, indicating that  
286 further confirmatory analyses should be performed. These results are used to create the  
287 corresponding validation contingencies of the classification model, as shown in Fig. 5b.

288

*Fig. 5*

289

290 **▪ PLS-DA**

291 The PLS-DA model was built using 4 latent variables, which explained 78.3% and 44.1%  
292 of the cumulative variance of both X- and Y-variable blocks, respectively. A threshold  
293 value of 0.5 was established as a decision criterion for the classification of the samples;  
294 scores (weights)  $> 0.5$  correspond to TB and  $< 0.5$  to TBM, as can be observed in the  
295 classification plot represented by Fig. 6a. The validation contingencies of the PLS-DA  
296 classification model are shown in Fig. 6b. Note that all validation samples were correctly  
297 classified, even though some samples from the training set were misclassified. This  
298 demonstrates the powerful generalization capabilities of the PLS-DA model.

299

*Fig. 6*

300

301 **▪ SVM**

302 Support vectors machine (SVM) was performed using the radial basis function (RBF)  
303 kernel algorithm with the gamma and cost values studied in the  $10^{-6}$ - $10^{-10}$  and  $10^{-3}$ - $10^2$

304 ranges, respectively, and PLS compression with 4 LVs. The classification results for both  
305 the training and validation samples are displayed in Fig. 7a. The results are almost the  
306 same as the PLS-DA ones, suggesting that sample TB70 should undergo further  
307 confirmatory analyses, since it is very close to the threshold value. The SVM validation  
308 contingencies are displayed in Fig. 7b.

309

*Fig. 7*

310

311 As a matter of comparison, the classification performance metrics for the classification  
312 models were calculated from the results of the validation contingencies (see Table 1) [22],  
313 considering TB as the target class. The most popular metrics are discussed here; however, the  
314 detailed explanation of each of them is out of the scope of this work and interested readers are  
315 kindly forwarded to ref [22] for specific details on this topic.

316

*Table 1*

317

318 In principle, satisfactory classifications lead to classification performance metrics close to 1  
319 and bad models to 0. For instance, Table 1 shows that PLS-DA and SVM models have a  
320 sensitivity (SENS) = 1, whilst 1iC-SIMCA a and b yields SENS = 0.83, which indicates that  
321 PLS-DA and SVM models classify better the TB samples than 1iC-SIMCA. Specificity  
322 (SPEC) indicates that the TBM samples are correctly classified, being better for PLS-DA and  
323 SVM models with a value = 1 than for 1iC-SIMCA a and b with SPEC = 0.50 and 0.33,  
324 respectively. In fact, the 1iC-SIMCA b model, validated with all the TBM samples, provided  
325 worse classification results than 1iC-SIMCA a, validated with fewer TBM samples.

326 Additionally, the positive predictive value (PPV) (so-called precision) informs on the  
327 proportion of agreements in relation to all assigned values of TB class whilst the negative  
328 prediction value (NPV) takes into account the ratio between agreements and the total number  
329 of TBM samples. For PLS-DA and SVM those metrics were = 1, whereas for the 1iC-SIMCA  
330 a and b models PPV were = 0.71 and 0.26, and NPV = 0.67 and 0.88, respectively. The  
331 overall classification rate (OCR) was 100%, 100% and 83% for PLS-DA, SVM and 1iC-  
332 SIMCA, respectively, and the Matthews correlation coefficient (MCC) –which might be  
333 considered a compendium of the overall classification ability of the models– was 1.0, 1.0 and  
334 0.36 for the same classification models.

335 When the validation set 'a' is applied on the 1iC-SIMCA model, the validation results are  
336 relatively good; however, the results are fictitious as this set does not represent the reality of  
337 the sample population. The good results are due to the fact that in the validation set 'a' only 4  
338 TBM samples (non-target class) are considered, but when the number of TBM samples is  
339 increased (validation set 'b'), the model does not classify well. That is, the model classifies  
340 almost all TBM samples as belonging to the TB class, which is related to the results shown in  
341 the exploratory analysis and the no clustering tendency of the classes, so it is not possible to  
342 establish regions for each of them. Therefore, the SIMCA class modelling method is not  
343 suitable for the purpose of this study.

344 The classification ability of the models obtained in this study (PLS-DA and SVM models) are  
345 better than others previously reported for different purposes (despite a direct, straightforward  
346 comparison is not possible) applying PCA-linear discriminant analysis (LDA), with an overall  
347 classification rate (OCR) of 90.02%, SENS = 0.90 and SPEC = 0.96 [17]. Furthermore, in a  
348 previous study [18] in which nine models were built using mean-centered UV-Vis  
349 spectroscopic data to differentiate various classes of Tequila, it was found that nonlinear  
350 models behaved better than linear ones (EFFIC > 0.94).

351 In this context, it is worth noting that class modeling methods, such as 1iC-SIMCA, are  
352 particularly suitable for real-world authentication problems where the target class is always  
353 defined from the authentic or genuine product and is modeled with a large number of samples,  
354 since it is less common to find adulterated samples. This approach has a great potential when  
355 the ideal scenario with sufficient number of authentic samples (target class) are available,  
356 being capable to properly identify new samples obtained from non-authentic products and  
357 differentiate them from those specimens of genuine ones. However, for this particular study,  
358 the available samples to build a more reliable 1iC-SIMCA model were limited, since Tequila  
359 Blanco 100% agave is only produced in certain regions of México and the accessibility of a  
360 variety of samples is rather narrow. A good alternative to address this situation is the use of  
361 discriminant methods, such as PLS-DA and SVM, particularly in this study, because it aimed  
362 at classifying two mutually excluding classes ('100 agave' and 'mixto') of the same quality sort  
363 of tequila ('Tequila Blanco'). In fact, it was evidenced that the validation results of the 1iC-  
364 SIMCA model depend on the number and type of samples included in the test set, but PLS-  
365 DA and SVM models provided better ability to correctly classify samples from both classes.  
366 However, this discriminant strategy is not free from the drawback of misclassifying new  
367 samples coming from non-genuine products with some different composition from those  
368 already used in the training step, which is a risk that practitioners must evaluate and take into  
369 account when extending the application of the method.

### 370 ***3.5. Alcoholic content quantitation***

371 A PLSR-based quantitation analytical method was calibrated to predict the alcoholic content  
372 of the Tequila samples. As detailed above, the reference values were obtained by the CRT  
373 following the official method. The PLSR model was built using mean centering to preprocess  
374 the spectra and including 5 LVs in the model which explained 73.6 and 97.1% of the  
375 cumulative variance for the X- and Y-variable blocks, respectively. Fig. 8 compares the PLSR



376 predicted alcoholic contents against the total alcoholic content reported by the CRT. The  
377 evaluation of this model was performed with the quantitation performance metrics, as  
378 observed in Table 2.

379

*Fig. 8*

380

*Table 2*

381

382 The first quantitation performance metric is the coefficient of determination ( $R^2$ ) with a value  
383 = 0.971, evidencing a good fitting. The following four metrics are related to different sorts of  
384 errors the model might present (root mean square error, mean absolute error, median absolute  
385 error and standard error of validation), all of them with values less than 4%; the sixth metric is  
386 the standard deviation of validation residuals ( $SDV = 2.7\%$ ), indicating that the agreement of  
387 the predictions of the empirical model with the reference values is high, which results in a  
388 quite good predictive ability.

389 Note that PLSR has been previously applied to predict the alcoholic content of different  
390 Tequilas using FTIR, obtaining very good results [19]. Moreover, a vector network analyzer  
391 with an open-ended coaxial probe kit was used for the same purpose [31].

392 PLSR has also been applied to quantitate the furfural, 2-acetylfuran and 5-methylfurfural  
393 content in White Tequilas and Mezcal samples with acceptable results [29]. It would have  
394 been interesting to compare the results obtained here with those of another report in which  
395 SORS was applied to study the adulteration of Vodka, Gin and Whisky with methanol, but  
396 prediction of the alcoholic content was not considered [10].

397

398 **4. Conclusions**

399 Economic losses for the industry of alcoholic beverages and societal health problems are two  
400 relevant consequences of the adulteration and counterfeiting of commercialized spirits, which  
401 have not ceased over the years. To streamline the authentication surveillance of these  
402 products, current official rearguard methods need to be complemented with vanguard, faster  
403 and reliable *in-situ* screening analytical methods. In this regard, the present study reports for  
404 the first time the combination of the SORS analytical technique and chemometrics to  
405 discriminate between 100% agave and mixed White Tequilas and to predict their alcoholic  
406 content. It should be noted that the potential of the *in-situ* non-invasive SORS measurement  
407 implemented here has been verified by means of a similarity analysis. This demonstrated that  
408 the spectra obtained after analyzing Tequilas through the original bottle and through amber  
409 vials are almost the same, obtaining nearness indexes close to 1. Afterwards, models were  
410 developed and assessed with several classification performance metrics, which indicated that  
411 satisfactory classifications and predictions were achieved. PLS-DA and SVM presented the  
412 best OCR = 100%, evidencing that the combination of SORS and some chemometric methods  
413 is able to discern among 100% agave and mixed White Tequilas. Finally, a PLSR quantitation  
414 model demonstrated an excellent ability to predict the alcoholic content of the samples.

415 The approach presented here offers an alternative analytical method for routine authentication  
416 tasks undergone by official regulatory bodies. It is reliable and fast for *in-situ* screening  
417 purposes and, can complement and accelerate the quality control and authentication processes  
418 of commercial spirits, such as Tequila.

419

420 ***Conflicts of interest***

421 The authors declare that they have no conflict of interest.

422

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431

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Table 1. Summary of classification performance metrics for liC-SIMCA, PLS-DA and SVM models.

Metrics	liC-SIMCA		PLS-DA	SVM
	a	b		
	<i>Target class (100% agave White Tequila, TB)</i>			
Sensitivity (SENS)	0.83	0.83	1.00	1.00
Specificity (SPEC)	0.50	0.33	1.00	1.00
False positive rate (FPR)	0.50	0.67	0.00	0.00
False negative rate (FNR)	0.17	0.17	0.00	0.00
Positive predictive value (PPV) (precision)	0.71	0.26	1.00	1.00
Negative predictive value (NPV)	0.67	0.88	1.00	1.00
Youden index (YOUD)	0.33	0.17	1.00	1.00
Positive likelihood rate (LR(+))	1.67	1.25	–	–
Negative likelihood rate (LR(-))	0.33	0.50	0.00	0.00
Classification odds ratio (COR)	5.00	2.50	–	–
F-measure (F)	0.77	0.40	1.00	1.00
Discriminant power (DP)	0.39	0.22	–	–
Efficiency (or accuracy) (EFFIC)	0.70	0.44	1.00	1.00
Misclassification rate (MR)	0.30	0.56	0.00	0.00
AUC (correctly classified rate) (CCR)	0.67	0.58	1.00	1.00
Gini coefficient (Gini)	0.33	0.17	1.00	1.00
G-mean (GM)	0.65	0.53	1.00	1.00
Matthews' correlation coefficient (MCC)	0.36	0.15	1.00	1.00
Chance agreement rate (CAR)	0.54	0.39	0.52	0.52
Chance error rate (CER)	0.48	0.35	0.48	0.48
Kappa coefficient (KAPPA)	0.35	0.09	1.00	1.00
PROB (TB/TB)	0.71	0.26	1.00	1.00
PROB (nTB/nTB)	0.67	0.88	1.00	1.00
PROB (TB/nTB)	0.33	0.13	0.00	0.00
PROB (nTB/TB)	0.29	0.74	0.00	0.00

The hyphen "-" signifies that the performance feature cannot be determined since it involves a division between zero.

a and b: models validated using 10 (6 TB and 4 TBM) and 27 (6 TB and 21 TBM) samples as external validation sets, respectively.

Table 2. Performance metrics in the quantitation of the alcoholic content of the Tequila samples that constitute the validation set.

<b>Metrics</b>	<b>Value (%)</b>
Coefficient of determination ( $R^2$ )	0.971
Root mean square error (RMSE)	3.32
Mean absolute error (MAE)	1.82
Median absolute error (MdAE)	2.61
Standard error of validation (SEV)	3.14
Standard deviation of validation residuals (SDV)	2.65

## Figure legends

**Figure 1.** Raman spectra of a '100% agave' White Tequila sample (TB) and a 'mixed' White Tequila (TBM) one.

**Figure 2.** Similarity plots of four sample pairs of White Tequila (S1-S4) measured through the original bottle (BS) and amber vial (VS), considered as the reference spectrum.

**Figure 3.** Exploratory PC1 vs PC2 scores plot from the 51 samples PCA model showing two different categories of White Tequilas. TB: 100% agave White Tequila (n=30) and TBM: mixed White Tequilas (n=21).

**Figure 4.** Exploratory LV2 vs LV3 scores plot from the 51 samples PLS model showing two different categories of White Tequilas. TB: 100% agave White Tequila (n=30) and TBM: mixed White Tequilas (n=21).

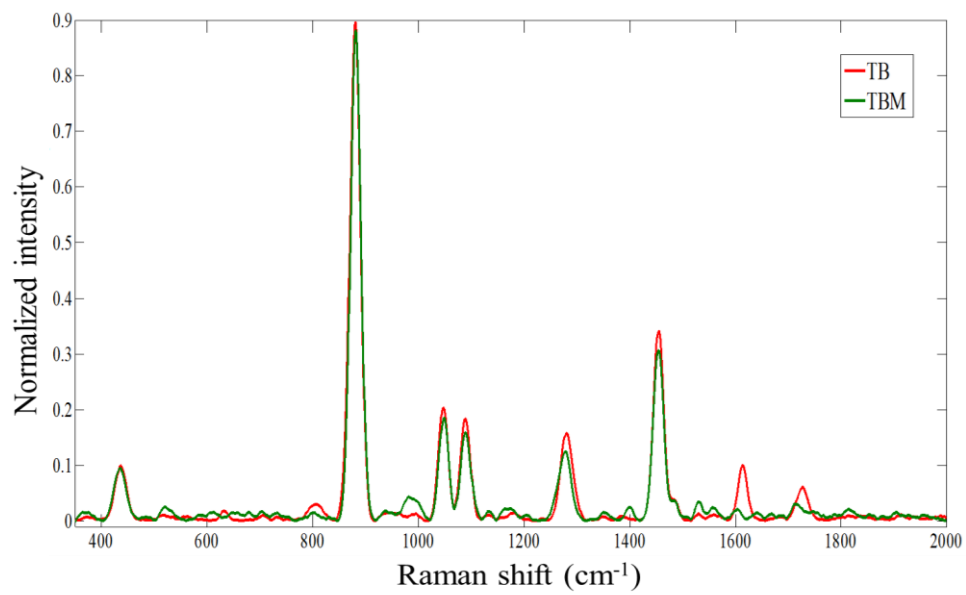
**Figure 5.** (a) Classification plot (a) and (b) validation contingencies for the one input-class SIMCA classification model. Class 1: target class (TB: '100% agave' Tequila); class 2: non-target class (TBM: 'mixed Tequila') (The magenta-marked samples in figure 5a are the misclassified samples).

**Figure 6.** (a) Classification plot and (b) validation contingencies for the PLS-DA classification model. Class 1: target class (TB: '100% agave' Tequila); class 2: non-target class (TBM: 'mixed Tequila'). (The dashed line in figure 6a indicates the 0.5 threshold level).

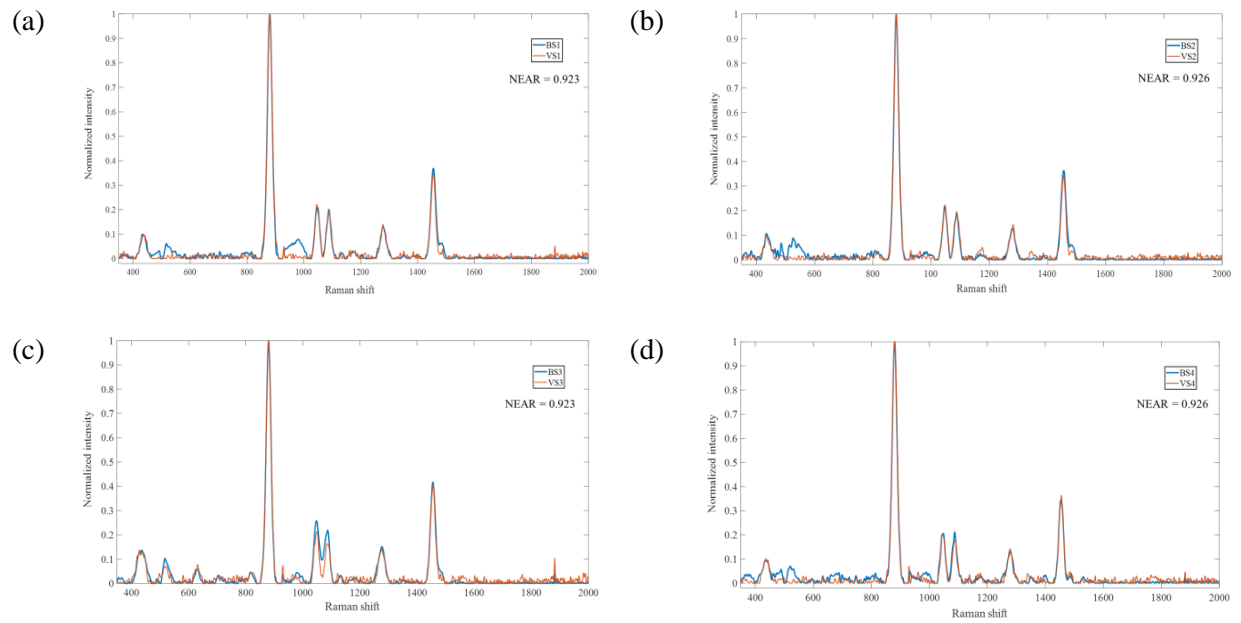
**Figure 7.** (a) Classification plot and (b) validation contingencies for the SVM classification model. Class 1: target class (TB: '100% agave' Tequila); class 2: non-target class (TBM: 'mixed Tequila'). (The dashed line in figure 7a marks the 0.5 threshold level).

**Figure 8.** PLSR alcoholic predictions (% v/v) for White Tequila samples. (a) Calibration curve, and (b) alcoholic content plot of the validation set samples. The circles are colored according to the predicted alcoholic content from the vertical color scale. Each sample displays the predicted value against the real value of alcoholic content, which is underlined.

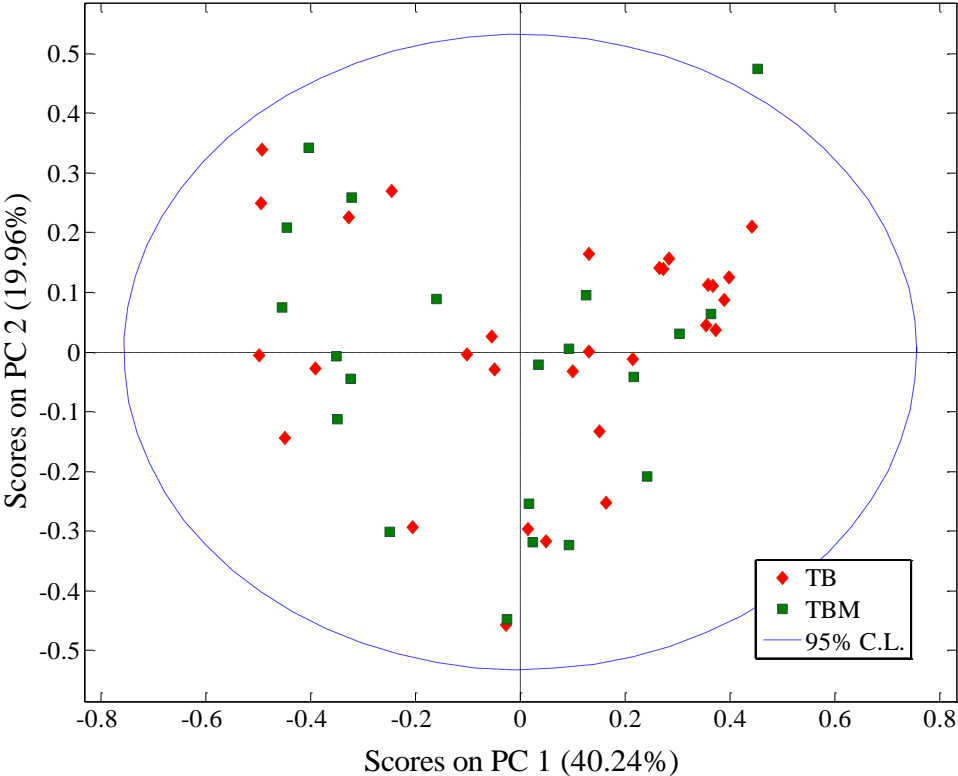
<Figure 1>



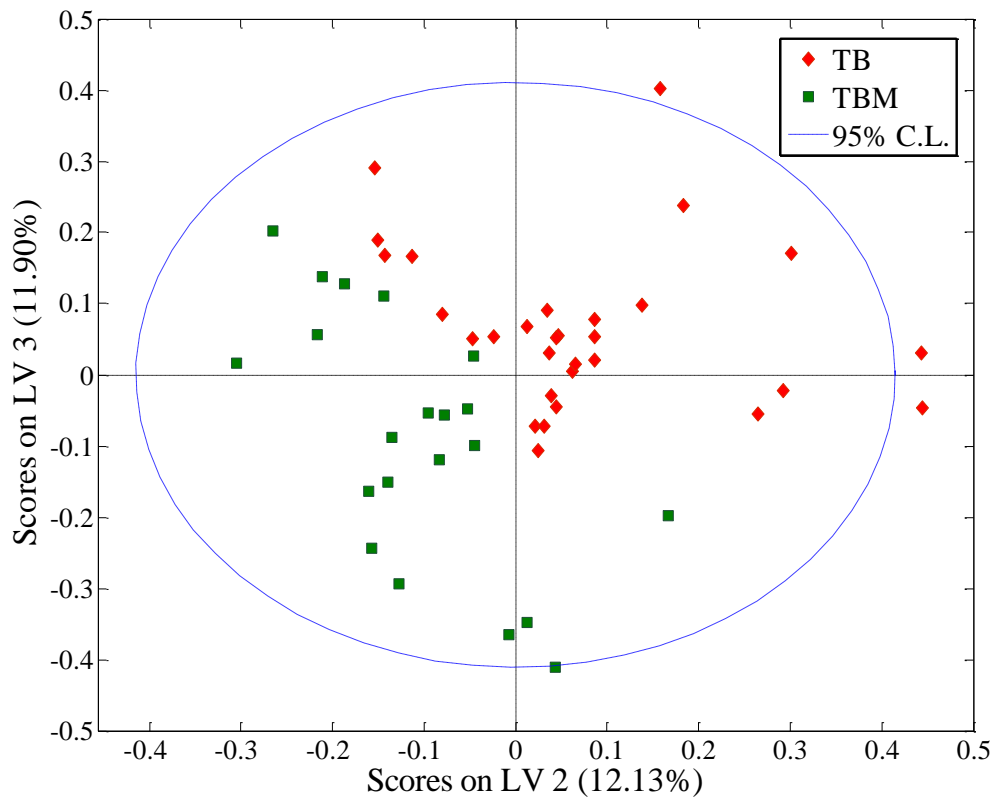
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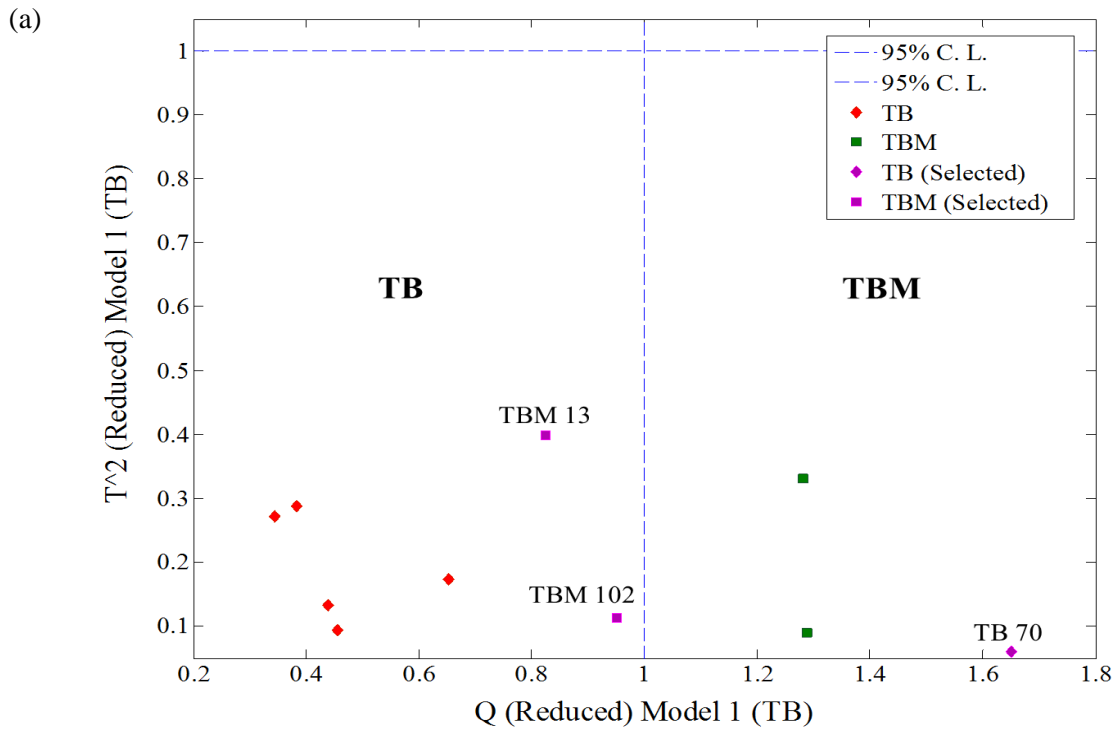
<Figure 3>



<Figure 4>



<Figure 5>

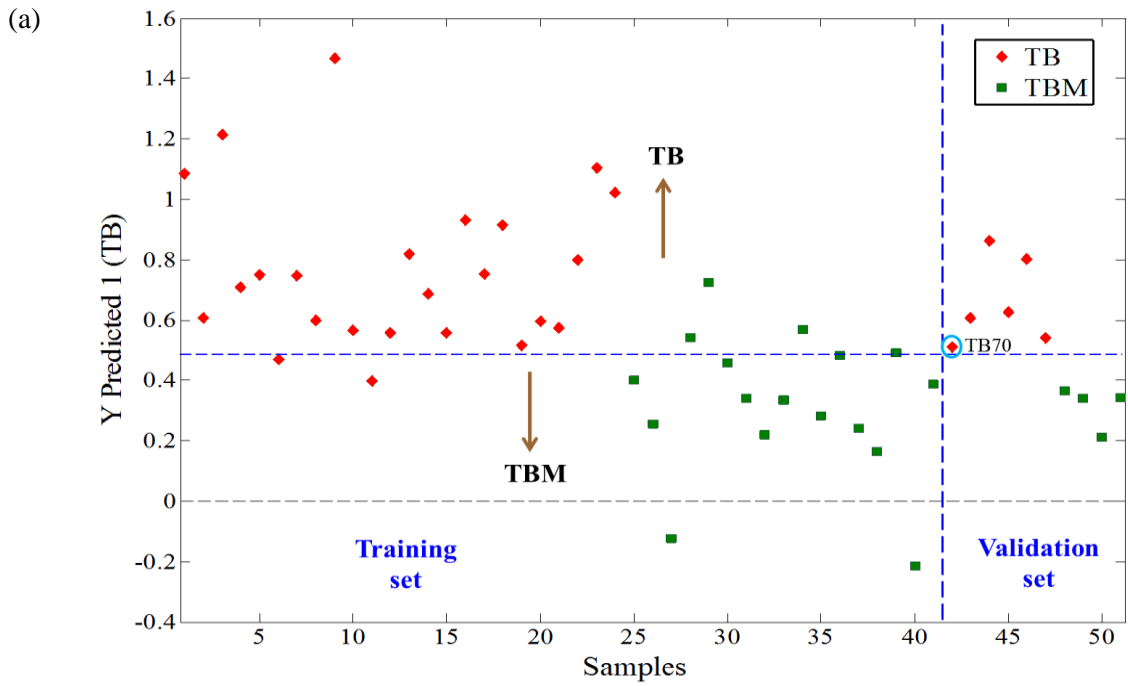


(b)

		6	4	10
Assignment	Inconclusive (I)	0	0	0
	Class 2 (TBM)	1 (8.33%)	2 (25%)	3
	Class 1 (TB)	5 (41.67%)	2 (25%)	7
		Class 1	Class 2	
		Actual		



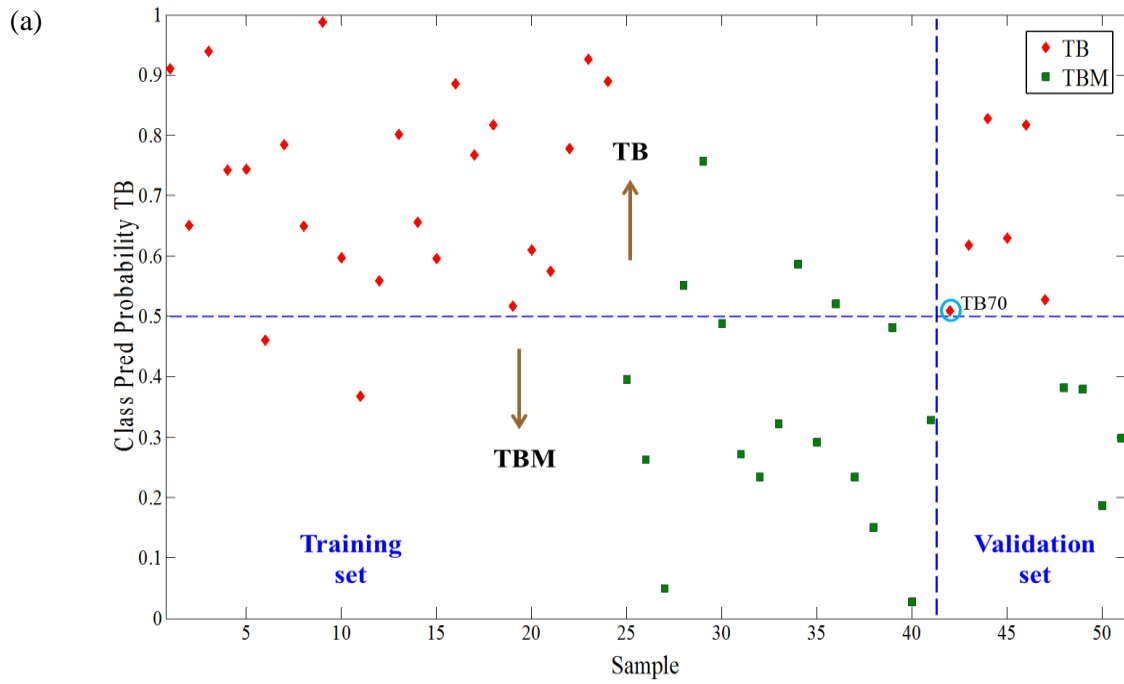
<Figure 6>



(b)

		<b>6</b>	<b>4</b>	<b>10</b>
<b>Assignment</b>	<b>Inconclusive (I)</b>	0	0	<b>0</b>
	<b>Class 2 (TBM)</b>	0	4 (50%)	<b>4</b>
	<b>Class 1 (TB)</b>	6 (50%)	0	<b>6</b>
		<b>Class 1</b>	<b>Class 2</b>	
		<b>Actual</b>		

<Figure 7>



(b)

		<b>6</b>	<b>4</b>	<b>10</b>
<b>Assignment</b>	<b>Inconclusive (I)</b>	0	0	<b>0</b>
	<b>Class 2 (TBM)</b>	0	4 (50%)	<b>4</b>
	<b>Class 1 (TB)</b>	6 (50%)	0	<b>6</b>
		<b>Class 1</b>	<b>Class 2</b>	<b>Actual</b>

<Figure 8>

