1	Sampling effects on the quantification of sodium content in infant formula using laser
2	induced breakdown spectroscopy (LIBS)
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18 Abstract

19 In the present work, laser-induced breakdown spectroscopy (LIBS) was employed to predict 20 the sodium content of infant formula (IF) over the range of 0.5–4 mg Na/g. Calibration 21 models were built using partial least squares regression (PLS), correlating the LIBS spectral 22 data with reference Na contents quantified by atomic absorption spectroscopy (AAS). The 23 aim of this study was to demonstrate the ability of LIBS as a rapid tool for quantifying 24 sodium in IF, but also to explore strategies concerning the acquisition of measurements with 25 LIBS. A range of different pre-processing techniques, measuring depths (repetition of laser 26 shots) and accumulations were evaluated in terms of PLS performance. The best calibration 27 model was developed using the third-layer spectra normalised by the H I 656.29 nm emission 28 line, yielding a coefficient of determination (\mathbb{R}^2) of 0.93, and root-mean-square errors 29 (RMSE) of 0.37 and 0.13 mg/g for cross-validation and validation, respectively.

30

31 Industrial relevance

Improving productivity and robustness of manufacturing processes, yet satisfying increasing concerns and strict regulations on the quality and safety of infant products could be achieved through the introduction of optical analytical techniques with real-time capabilities during processing. In this paper, LIBS is proposed as a potential cost-effective screening tool that can provide fast elemental composition analysis of IF. Specifically, the application of LIBS and multivariate data analysis for predicting sodium content over a range in conformity with regulatory guidelines is discussed in this work.

39

40 Keywords

41 LIBS; Infant formula; Sodium; Partial least squares regression; sampling

42 **1. Introduction**

43 Infant formula (IF) is an industrially produced food intended as a substitute for breast milk. 44 IFs are typically based on cow's milk, and followed by several adjustments and addition of 45 ingredients in order to bring the composition closer to that of human milk (Blanchard, Zhu, & Schuck, 2013). Infancy is a crucial period of growth and development, hence IF's 46 47 composition (e.g. fat, proteins, minerals, etc.) and manufacturing practices are strictly 48 regulated by national authorities to ensure the safety and nutrient profile of infant formula 49 products (Jiang, 2014; Montagne, Van Dael, Skanderby, & Hugelshofer, 2009). 50 Sodium is an essential mineral; it is the main cation in extracellular fluid playing a vital role 51 in the regulation of osmolarity, acid-base equilibrium, active transport across cells and 52 membrane potential (Guo, 2014). Although a minimum intake is indispensable for healthy 53 functioning, an excessive consumption of sodium in the human diet is related to higher blood 54 pressure and an increased risk of developing cardiovascular diseases (Masotti, Erba, De Noni, 55 & Pellegrino, 2012; Tamm, Bolumar, Bajovic, & Toepfl, 2016). With regard to infancy, 56 studies have also associated an excessive sodium intake with increased blood pressure in the 57 later stages of life, indicating that blood pressure may track with age (Campbell et al., 2014; 58 John et al., 2016). 59 Conventional well-established methods for mineral analysis in infant formula include atomic 60 absorption spectroscopy (AAS), inductively coupled plasma optical emission spectroscopy 61 (ICP-OES) and inductively coupled plasma mass spectroscopy (ICP-MS) (Poitevin, 2016). 62 These methods, despite their high sensitivity and accuracy, generally require time-consuming 63 and laborious sampling procedures and the use of chemical reagents such as acids and gases, 64 as well as an associated high cost of consumables (e.g. argon) (Wu & Sun, 2013). Laser-induced breakdown spectroscopy (LIBS) is an analytical technique based on optical 65 66 emission spectroscopy in which laser pulses are employed as the excitation source to

67 vaporise, atomise and ionise a small part of the target's material. As a result, plasma arising 68 from the sample surface is generated from which photons are released form the excited 69 species in the plasma returning to their ground state levels of energy which can be analysed 70 with spectrometers to infer the elemental composition of the sample (Cremers & Radziemski, 71 2013). LIBS, yet recent in the area of food analysis, has gained remarkable popularity in the 72 last few years with an increase in the number of publications and extensive reviews 73 concerning food samples (Maria Markiewicz-Keszycka et al., 2017; Sezer, Bilge, & Boyaci, 74 2017). The advantages that LIBS offers compared to the conventional methods are its speed, 75 a relatively low cost, little to no sample preparation and elemental surface mapping 76 capabilities (Dixit et al., 2017; Kim, Kwak, Choi, & Park, 2012). Further attractive features 77 include: remote sensing, as it constitutes an entirely optical technique, and suitability for on-78 /at-line applications, altogether allowing the technology to be considered a potential process 79 analytical technology (PAT) for qualitative and quantitative chemical analysis (Cullen, 80 Bakalis, & Sullivan, 2017) (for PAT literature the reader is referred to: Misra et al., 2015; van 81 den Berg et al., 2013). Nonetheless, LIBS also has limitations or drawbacks, especially 82 concerning quantitative analyses. Some of these limitations include signal fluctuations on a 83 shot-to-shot basis (Tognoni & Cristoforetti, 2016) and difficulties in establishing good 84 calibration curves due to strong matrix effects (Ferreira et al., 2010; Lei et al., 2011). Several 85 publications evaluating and discussing strategies with the goal of overcoming such problems 86 can be found in the literature (dos Santos Augusto, Barsanelli, Pereira, & Pereira-Filho, 2017; 87 El Haddad, Canioni, & Bousquet, 2014; Jantzi et al., 2016). 88 In this study, LIBS and multivariate data analysis with partial least squares regression (PLS) 89 was employed to predict the sodium content of IF samples. In order to provide for reference 90 Na contents, atomic absorption spectroscopy (AAS) was used. The aim of this study was to

91 demonstrate the ability of LIBS as a rapid screening tool for quantifying sodium over a range

92 relevant to IF manufacturing, offering a means for industries to rapidly verify target mineral
93 contents. Furthermore, strategies concerning the acquisition of measurements with LIBS were
94 explored, namely the repetition of laser shots on a single location. Such an approach
95 examines the impact of measuring the inner layers of the sample and, whether to accumulate
96 laser shots or use the spectra collected from a single layer.

97 2. Material and methods

98 2.1. Sample preparation

99 Commercial powdered IF and follow-on formulas (formulas intended for children over 6
100 months of age) were acquired from a local supermarket in Dublin, Ireland. Lactose (α-lactose
101 monohydrate ≥ 99 %) and sodium chloride (NaCl ≥ 99 %) were purchased from Sigma
102 Aldrich (St. Louis, MO, USA).

103 Samples with varying content of sodium were prepared by blending IF with sodium chloride 104 or lactose, whether the goal was to increase or decrease the sodium content in the mix. In 105 total, 7 samples were obtained, including one sample which consisted only of IF. The 106 selected range of sodium was approx. from 0.5 to 4 mg/g (concentrations corresponding to 107 the lowest and highest Na content samples, respectively). This range was intended to cover 108 the regulatory sodium levels provided by the Codex Alimentarius Commission (Codex, 109 2007). Constituents of the mixtures (IF, NaCl and lactose) and follow-on formulas were 110 ground and pre-mixed using a laboratory blender (8011G, Waring Laboratory Science, CT, 111 USA) equipped with rotatory stainless-steel blades for 2 minutes to ensure there were no 112 aggregates occurring in the powders, with the goal of improving subsequent blending 113 performance. Dry mixing was then carried out using a laboratory V-mixer (FTLMV-1L&, 114 Filtra Vibracion S.L., Spain) for 20 minutes. In order to ensure reproducibility, two independent batches were prepared (batch 1 and batch 2). Each batch was composed of the 115 116 aforementioned 7 samples divided into: 5 calibration samples (referred to as C1-C5),

employed for PLS modelling, and 2 validation samples (V1, V2) to test the robustness of the models. In addition to these validation samples, 2 different follow-on formula brand samples (V3, V4) were used to assess the ability of the calibrations for predicting mineral content in infant products with different formulations.

121 For LIBS analysis, samples were pelletized by pressing approx. 400 mg of each sample using

a manual hydraulic press fitted with a 13 mm pellet die (Specac Ltd., UK) at 10 tons for 3

123 minutes. Pellets were prepared in triplicates (3 replicates per sample), giving a total number

124 of 48 pellets. The two batches of samples were measured on different days.

125 2.2. Atomic absorption spectroscopy (AAS)

126 AAS was selected as the reference method for sodium quantification in IF mixtures. Na

127 contents were established using a Varian 55B AA spectrometer (Varian, United States)

128 following the standard method 985.35 for mineral determination in IF of the AOAC (Official

129 Methods of Analysis of AOAC International) with slight modifications. Approximately 1.5 g

130 of each sample was transferred to a crucible in triplicates (3 replicates). Crucibles were

131 placed on a hot plate and heated until smoking ceased. Organic matter was then decomposed

132 by dry ashing in a muffle furnace at 525°C for 4 h. Ashes were dissolved in 50 mL 1 M nitric

133 acid. A further dilution step was required to bring concentrations within the linear range of

134 the instrument (0–1 ppm).

135 Calibration curves were established by using aqueous standards prepared from a commercial

136 sodium stock solution (Sodium standard for AAS – 1,000 mg/L, Sigma-Aldrich). Sodium

137 absorbance was measured at 589 nm with a slit width of 0.5 nm. All replicates and batches

138 were measured on different days.

139 2.3. LIBS instrumentation and measurements

140 2.3.1. Instrument set-up

141 LIBS spectra were recorded using a LIBSCAN-150 system (Applied Photonics Ltd, UK)

142 described in a previous publication (X. Cama-Moncunill et al., 2017). The system was fitted

- 143 with a 150 mJ Q-switched Nd:YAG laser (Ultra, Quantel laser, MT, USA) operating at 1064
- nm and a pulse duration of 5 ns, coupled to six fibre-optic spectrophotometers (AvaSpec,

145 Avantes spectrometers, Netherlands) covering the wavelength range of 181–904 nm.

146 Moreover, the system was equipped with a miniature CCD camera which enabled the

- 147 monitoring of the measurements.
- 148 For the experiments, plasma emission was analysed with a delay time of 1.27 µs and an

149 integration time of 1.1 ms. The laser was operated with a firing repetition rate of 1 Hz.

150 2.3.2. Sampling method

151 Pellets were measured individually using a sample chamber equipped with a three-axis

152 translation stage (Applied Photonics Ltd, UK) which facilitated the acquisition of spectra at

153 multiple locations of the pellet surface, that is, 100 locations following a 10×10 grid pattern.

154 Spectral acquisition was carried out by recording 5 consecutive laser shots (depth

155 measurements) at each of the 100 locations, giving a total number of 500 measurements per

156 pellet. Data resulting from these consecutive laser shots can be considered as spectra

157 corresponding to 5 different layers of the pellets, i.e. the repetitive firing of the laser at the

same location causes the ablation of the outer material penetrating and allowing to measure

159 deeper into the sample (Cremers & Radziemski, 2013).

160 Spectral data collected from the 5 laser shots were stored separately in order to assess the best

161 layer from which to build the sodium quantification model, and to allow subsequent

162 comparison between accumulated and non-accumulated laser shots.

163 *2.4. Data analysis*

Data analysis was performed with R (R Core Team, 2014) using the R package "pls" (Mevik,
Wehrens, & Liland, 2015) for conducting PLS (partial least squares regression), as well as
other in-house functions.

Firstly, the average of the LIBS spectra collected at multiple locations was calculated for 167 168 each layer, resulting in 5 spectra per pellet. Data was then divided into a training data set (N=30) and a test set (N=12), additionally the follow-on formula extra validation samples 169 170 (N=6) were tested. Prior to PLS modelling, combinations of different pre-processing 171 techniques and normalisation methods were applied to the spectra with the aim of reducing 172 the signal fluctuations due to extraneous sources of variability and to minimize any matrix 173 effects (Sobron, Wang, & Sobron, 2012). Specifically, the techniques explored were: baseline 174 correction (R package "baseline"), second derivative and standard normal variate (SNV). 175 Spectral normalisation using other approaches, including normalisation by an internal 176 standard and the Euclidean norm, were also explored. 177 PLS calibration models using the different pre-processing techniques were developed for 178 each of the 5 layers of the pellets. The performance of each model was evaluated by the 179 leave-one-out root-mean-square error of cross-validation (RMSECV) technique, as well as 180 the root-mean-square error of prediction (RMSEP). The wavelength range used for the 181 modelling was limited to 560–825 nm since this region encompassed the main Na emission 182 lines, while decreasing the total number of variables that do not contain useful peaks 183 (Moncayo, Manzoor, Rosales, Anzano, & Caceres, 2017). 184 In order to provide for a comparison between the accumulated and non-accumulated shots, 185 spectra corresponding to the different layers were summated to one another so that 2, 3, 4 and

186 5 accumulations were obtained. PLS modelling of the accumulated spectra was then carried

187 out, and their resulting performances were compared to those of the single-layer-spectra188 models.

189 The limit of detection was computed according to the pseudounivariate approach (LOD_{pu}) for

190 PLS models as proposed in a publication elsewhere (Allegrini & Olivieri, 2014) in

191 accordance with IUPAC official recommendations. LOD_{pu} calculation was performed as

shown in Eq. 1.

193
$$LOD_{pu} = \frac{3.3}{S_{pu}} \left[\left(1 + h_{0 \min} + \frac{1}{I} \right) var_{pu} \right]^{1/2}$$
 (1)

where S_{pu} is the slope of the pseudounivariate line, $h_{o \min}$ is the minimum leverage when the analyte concentration is zero, *I* the number of samples employed for calibration, and var_{pu} is the variance of the regression residuals.

197 **3. Results and discussion**

198 *3.1. AAS*

In AAS, the accuracy of the results relies heavily upon the calibration curve established from reference standard solutions of the desired element. Good calibration curves were obtained rendering values for the coefficient of determination (R^2) \geq 0.99. Sodium contents of the IF samples determined with AAS, expressed in mg/g, are shown in Table 1.

203 3.2. LIBS spectral features

204 An initial exploratory analysis of the LIBS spectra was conducted in order to determine the

205 principal differences among the samples studied. For comparison purposes, the averaged

206 spectra of pellets corresponding to the lactose-IF mixture (C1, approx. 0.5 mg Na/g), pure IF

207 (C2, approx. 1.3 mg Na/g) and the sodium chloride-IF mixture (C5, approx. 3.7 mg Na/g) are

- shown in Fig. 1. In the figure, several of the most important spectral lines of elements
- 209 occurring in the spectra can be seen. The main element emission lines in the spectra were
- 210 identified using the NIST database (Kramida, Ralchenko, Reader, & NIST ASD team, 2016).
- 211 These emission lines included: C I 247.86 nm, Ca II 393.37; 396.85 nm, Ca I 422.67; 558.88;

212 612.22; 616.22 nm, H I 656.29 nm, N I 744.23; 746.83 nm, K I 766.49; 769.90 nm, O I

213 777.19 nm and Na I 589.05; 589.59 nm. Moreover, three Na I lines were identified at 568.26,

214 568.82 and 819.48 nm. Other possible Na lines in the spectra were discarded and not

215 considered for quantitative analysis since the intensities at these wavelengths were marginal,

216 which is consistent with the NIST guidelines for sodium.

217 3.3. Multivariate analysis with PLS

218 PLS is a method for predicting a quantitative response (i.e. sodium content), stored in a

219 matrix Y, from numerous predictor variables (i.e. spectral data), stored in a matrix X. In order

to do so, it decomposes simultaneously the two matrices into new variables, known as factors

221 or latent variables (LV), in such a way that they explain as much as possible of the covariance

between X and Y. A multivariate linear model is then fitted using the latent variables to

223 predict the quantitative response (Abdi, 2010).

224 PLS modelling has been demonstrated to successfully develop quantitative calibration models

from LIBS spectral data of food samples in previous publications (Andersen, Frydenvang,

Henckel, & Rinnan, 2016; Bilge et al., 2016; M. Markiewicz-Keszycka et al., 2018). In the

227 present study, PLS was employed to build the calibration models for the determination of

sodium content by correlating the pre-processed LIBS spectra in the wavelength range of

229 560–825 nm to the reference Na contents extracted from AAS analysis.

230 *3.3.1. PLS modelling: performance of sampling methods and spectral pre-processing*

As previously mentioned, different techniques and normalisation methods were explored as

232 pre-processing techniques of the spectra prior to modelling. To this end, various calibrations

233 were developed using the approaches detailed in section 2.4. A summary of PLS

performances for these calibrations can be found in Table 2 (for briefness, this table only

includes some of the most relevant models). The criterion followed for establishing an

236 optimum number of LVs for each model considered a low value of RMSECV (root-mean-

square error of cross-validation) with a low number of LVs to avoid overfitting. In order to
determine the best calibration for quantifying sodium content in IF samples, both RMSECV
and RMSEP (root-mean-square error of prediction) were used.

240 With regards to pre-processing techniques, the best performances were obtained for 241 normalised spectra with SNV, Euclidean norm and normalisation using the H I at 656.29 nm 242 and Ca I at 422.6 nm emission lines as internal standards. All the methods above yielded 243 similar results for calibration (Table 2): e.g. the third-layer-spectra models (measurement 244 depth: 3) using these pre-processing techniques rendered values of almost 0.94 for the coefficient of determination (R²). These models also provided similar results for root-mean-245 246 square errors of cross-validation and prediction: third-layer-spectra models yielded values of 247 approx. 0.37 mg/g for RMSECV and values in the range of approx. 0.13-0.16 mg/g for 248 RMSEP. Other techniques such as baseline correction or normalisation with other internal 249 standards (C I at 247.9nm and K I at 766.4 nm) provided good calibrations and reasonable 250 validation performances. However, the RMSEP values were slightly higher than those 251 obtained with SNV, Euclidean, H I 656.29 nm or Ca I 422.6 nm. Second derivative preprocessing was found not to be effective for calibration showing low values of R^2 and R_{CV}^2 252 (coefficient of determination for cross-validation), as well as high values of root-mean-square 253 254 errors (RMSE, RMSECV).

Regarding the modelling of layers or depth measurements, it was observed that the third-layer spectra exhibited the best results regardless of the pre-processing techniques used. The first and second layers, while providing a good calibration, showed performances considerably lower for cross-validation and validation. The fourth and fifth layers exhibited an overall good performance, but with lower R^2 values and higher RMSECV and RMSEP as compared to the third layer. The effect of measuring deeper into the sample on spectral quality, and as a mechanism to avoid surface contamination has been previously investigated (R. Cama-

262 Moncunill et al., 2017). Similarly, in this publication PLS models were developed for 263 different layers of the samples with the aim of quantifying copper and iron contents in infant 264 formula premixes (blends used in IF manufacturing which are designed to contain specified 265 nutrients). The authors observed that PLS performances, especially with regard to validation, improved as the measuring depth increased. In the present study, this trend was also 266 267 observed, however, finding an optimum at the third measurement depth. It is worth noting 268 that depending on the laser energy and sample type, the optimum number of shots on the 269 same location may change substantially since these parameters affect the laser-material 270 interaction, for instance the size of the crater formed or the amount of ablated mass (Tognoni 271 & Cristoforetti, 2016).

272 Table 2 also shows the performances for some of the PLS models developed with the 273 accumulated spectra. In this regard, the modelling of accumulated spectra only proved to 274 yield notable better performances for the first two laser shots as compared to applying PLS 275 separately on these layers. A larger number of accumulations did not provide better models 276 than using the third-layer-spectra alone. In several publications, authors chose to accumulate 277 spectra as a means to mitigate signal fluctuations (Maria Markiewicz-Keszycka et al., 2017). 278 The fact that, in this work, accumulating spectra did not considerably improved the results 279 may be due to an already high sampling number (average of 100 locations) along with an 280 optimum of 3 laser shots, the first two of which ablate away the surface which may have been 281 contaminated.

Considering both pre-processing and sampling method, the best performing PLS model to
predict sodium content was the third-layer spectra which had been normalised using the H I
emission line at 656.29 nm.

285 *3.3.2.* Validation of the selected calibration model

The hydrogen-normalised third-layer-spectra model was used as the calibration to perform sodium content predictions. Fig. 2 (a) shows the values of RMSECV for each LV of this model. A number of 3 LVs was selected as further factors did not result in a notable improvement in terms of RMSECV while, at the same time, the quality of the predictions for the validation set decreased, indicating that a higher number of LVs could result in overfitting of the model. The first 3 main LVs explained approximately 95.7% of the total spectral variance.

Fig. 2 (b) shows the loading values for the first factor of the PLS model in the wavelength range assessed. One main sodium (Na I) emission line at 589.59 nm contributed to the loading values. Other Na I spectral lines were the doublet at 568.26 and 568.82 nm, and the emission line at 819.48 nm. These spectral lines had a relatively small contribution as compared to the sodium doublet at around 589 nm. Negative loading values were only observed for nitrogen (N I 744.23 and 746.83 nm) and oxygen (O I 777.19 nm), both elements showing minor values.

The PLS model exhibited an R^2 of 0.93 for the calibration. With regards to cross-validation, 300 an R_{CV}^2 value of 0.886 and an RMSECV of 0.373 mg/g were obtained, indicating a reasonable 301 302 fit and accuracy of the calibration. The validation of the PLS model was carried out by 303 predicting the Na contents of 2 samples not included in the training set with the aim of 304 evaluating the robustness of the model. The model exhibited a good prediction accuracy as indicated by a high R_p^2 (coefficient of determination for the validation set) of 0.967 and a 305 306 RMSEP value of 0.129 mg Na/g. Fig. 3 shows the PLS calibration curve with the predicted 307 values for the validation set. To further evaluate the closeness of the predictions to the actual 308 values of concentration, the relative error (RE) was calculated as reported elsewhere (Câmara 309 et al., 2017). The RE value of the validation set was 7.22%.

Additionally, Na contents for 2 follow-on formulas were also predicted in order to explore the model's response to different formulations of infant products. In this case, the predictions were not as accurate as the validation set, giving a RE value of 23.32%. However, this result may indicate that the model can provide reasonable predictions even with a certain degree of variability in the raw materials.

315 As mentioned before, the best performance was given by spectra collected after 3 laser shots. 316 To further investigate why the third layer provided better results, sodium content was 317 predicted, in this case, for each location in the 10×10 measuring grid. In order to do so, the 318 raw spectral data acquired from sample V2, chosen as a point close to the centre of the 319 calibration curve, was normalised by the hydrogen emission line without averaging the data 320 of multiple locations, i.e. obtaining 500 pre-processed spectra instead of 5. Na contents were 321 subsequently predicted employing the coefficients extracted from the PLS model. Fig. 4 322 shows a schematic representation of the V2 pellet displaying sodium content in each spatial 323 position for the first 3 measurement depths. The same intensity scale for the three 324 measurements was implemented to allow comparison. It can be observed that the predictions 325 for the third layer, Fig.4(c), provided a more homogeneously distributed sodium within the 326 analysed area.

327 The limit of detection of the model was estimated by following the pseudounivariate

328 approach as described in Eq. 2. The LOD value corresponding to the calibration model was

329 1.11 mg/g.

330 4. Conclusions

LIBS was successfully applied for quantifying sodium over a range in conformity with the product's regulatory guidelines, hence, demonstrating the feasibility of the technique as a potential screening tool for IF manufacturing. Multivariate analysis with PLS was applied to spectral data processed by a range of different pre-processing techniques, measuring depths

335 and accumulations. The resulting calibration models were compared in terms of PLS 336 performance: coefficients of determination and root-mean-square errors; for calibration (\mathbb{R}^2 , RMSEC), leave-one-out cross-validation (R_{CV}^2 , RMSECV) and validation (R_p^2 , RMSEP). The 337 338 best PLS calibration was obtained using the third-layer spectra normalised by the H I emission line at 656.29 nm, yielding a R^2 of 0.93 and a R^2_{CV} of 0.886. When performing 339 validation of this model, the resulting R_p^2 and RMSEP values were 0.967 and 0.129 mg Na/g 340 341 respectively, proving its ability to accurately predict samples not included in the calibration 342 set.

In this study, accumulation of the spectra on the same spot did not notably improve the performances of the PLS models as compared to using the third layer alone. Furthermore, chemical mapping with PLS of the analysed area (100 measurements in a 10×10 grid pattern) showed that sodium was more homogeneously distributed than for the first two layers. These results suggested that conditioning the surface of the pelletized sample, while keeping a low number of shots on the same spot, can provide a good predictive accuracy without the need of large sampling numbers.

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491 **Table 1**

492 Sodium contents in milligrams per gram of samples corresponding to calibration (C1–C5)

Sampla	e Constituents	Batch 1	Batch 2	Extra validation		
Sample	Constituents	Na content (mg/g) ^a	Na content (mg/g) ^a	Na content (mg/g) ^a		
C1	IF + lactose	0.48 ± 0.05	0.54 ± 0.03	_		
C2	IF	1.40 ± 0.21	1.34 ± 0.07	_		
C3	IF + NaCl	2.11 ± 0.11	2.07 ± 0.02	_		
C4	IF + NaCl	2.78 ± 0.16	2.72 ± 0.07	_		
C5	IF + NaCl	3.69 ± 0.54	3.74 ± 0.18	_		
V1	IF + lactose	0.93 ± 0.06	0.98 ± 0.06	_		
V2	IF + NaCl	2.22 ± 0.04	2.48 ± 0.21	-		
V3	follow-on	_	_	1.18 ± 0.04		
V4	follow-on	_	_	2.38 ± 0.36		

493 and validation (V1–V4) determined by AAS.

494 ^a Contents expressed as mean \pm standard deviation of three replicates.

495 **Table 2**

496 Summary of performances for the PLS models developed using different sampling methods

Experiment	Depth	Pre-	Calibration		Cross-validation		Validation		
		processing	LVs	\mathbb{R}^2	RMSEC	R_{CV}^2	RMSECV	R_p^2	RMSEP
Single layer	3	None	3	0.851	0.426	0.771	0.529	0.904	0.218
Single layer	1	H I 656.3	3	0.899	0.352	0.822	0.465	0.612	0.438
Single layer	2	H I 656.3	3	0.856	0.419	0.776	0.523	0.498	0.498
Single layer	3	H I 656.3	3	0.930	0.291	0.886	0.373	0.967	0.129
Single layer	4	H I 656.3	3	0.879	0.384	0.786	0.511	0.915	0.205
Single layer	5	H I 656.3	3	0.824	0.463	0.665	0.639	0.914	0.207
Accumulations	$4 (0^{a/4^{b}})$	H I 656.3	3	0.931	0.290	0.884	0.377	0.914	0.206
Accumulations	$5(0^{a}/5^{b})$	H I 656.3	3	0.916	0.320	0.856	0.419	0.935	0.179
Accumulations	$4(1^{a}/3^{b})$	H I 656.3	3	0.924	0.287	0.872	0.366	0.940	0.213
Single layer	3	Ca I 422.6	3	0.937	0.276	0.893	0.361	0.966	0.131
Single layer	3	C I 247.9	3	0.923	0.306	0.876	0.389	0.894	0.229
Single layer	3	K I 766.4	3	0.942	0.266	0.911	0.330	0.908	0.213
Single layer	3	SNV	3	0.938	0.276	0.888	0.369	0.945	0.164
Single layer	4	SNV	3	0.917	0.318	0.851	0.426	0.866	0.258
Single layer	5	SNV	2	0.865	0.405	0.816	0.473	0.911	0.210
Accumulations	$4 (0^{a/4^{b}})$	SNV	2	0.879	0.384	0.840	0.442	0.849	0.274
Accumulations	5 (0 ^a /5 ^b)	SNV	2	0.881	0.382	0.841	0.441	0.897	0.225
Accumulations	$4(1^{a}/3^{b})$	SNV	2	0.878	0.389	0.831	0.446	0.888	0.312
Single layer	3	Euclidean	3	0.938	0.274	0.889	0.367	0.950	0.157
Single layer	4	Euclidean	2	0.871	0.397	0.824	0.463	0.889	0.234
Single layer	5	Euclidean	2	0.865	0.405	0.816	0.474	0.910	0.211
Accumulations	4	Euclidean	2	0.879	0.383	0.839	0.443	0.811	0.306
Accumulations	5	Euclidean	2	0.881	0.381	0.841	0.441	0.874	0.250

497 and pre-processing techniques.

498 ^a Number of conditioning shots.

499 ^b Number of accumulated spectra.



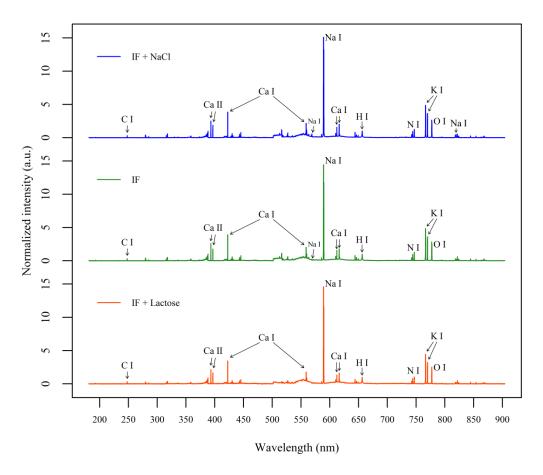
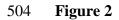
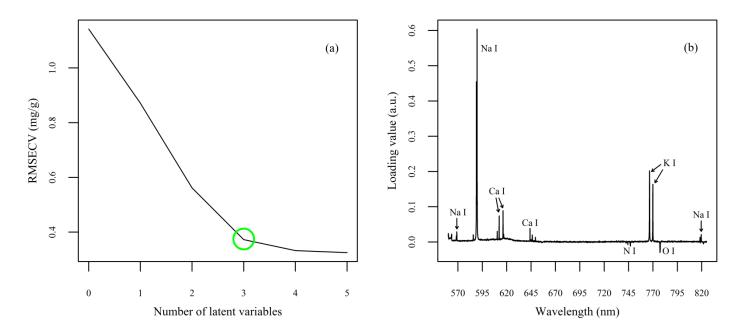


Fig. 1. Averaged spectra corresponding to, from top to bottom, the sodium chloride -IF mixture at approx. 3.7 mg Na/g, the pure IF sample at approx. 1.3 mg Na/g and the sodium lactose-IF mixture at approx. 0.5 mg Na/g. Spectra are vertically offset for illustration purposes.





505 Fig. 2. (a) RMSECV (root-mean-square error of cross-validation) for each number of PLS
506 factors or latent variables. (b) Loading values of each wavelength for the first latent variable.

Figure 3

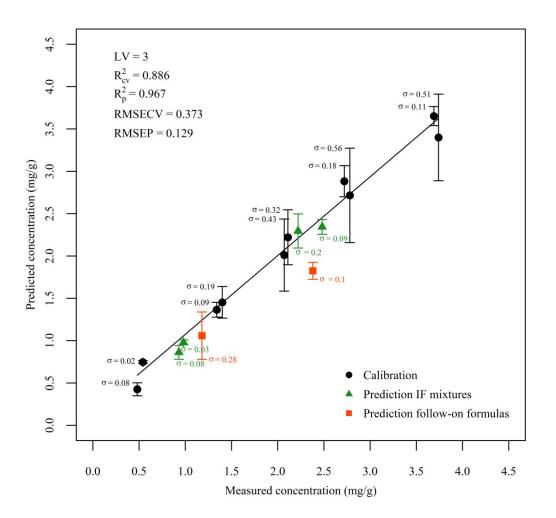
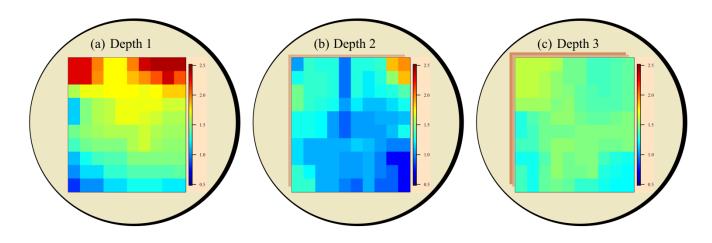


Fig. 3. PLS calibration model developed using the third-layer spectra and normalised by the
H I 656.29 emission line showing predicted Na contents for the validation and follow-on
formulas sets. Standard deviation values (σ) are expressed in mg/g.

511 Figure 4



512 **Fig.4.** Predicted sodium maps for the validation sample at 2.48 mg/g of sodium for the first

- 513 three depths: (a) first layer, (b) second layer, (c) third layer. The same intensity scale was
- 514 implemented for the three samples to facilitate comparison.