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## FEASIBILITY STUDY ON THE USE OF VISIBLE-NEAR INFRARED SPECTROSCOPY FOR THE SCREENING OF INDIVIDUAL AND TOTAL GLUCOSINOLATE CONTENTS IN BROCCOLI

José Miguel Hernández-Hierro, Juan Valverde, Salvador Villacreces, Kim Reilly, Michael Gaffney, Maria Lourdes González-Miret, Francisco José Heredia, and Gerard Downey

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**Feasibility Study on the Use of Visible-Near Infrared Spectroscopy for  
the Screening of Individual and Total Glucosinolate Contents in  
Broccoli**

José Miguel Hernández-Hierro<sup>†</sup>, Juan Valverde<sup>‡</sup>, Salvador Villacreces<sup>‡</sup>, Kim Reilly<sup>§</sup>,  
Michael Gaffney<sup>§</sup>, Maria Lourdes González-Miret<sup>†</sup>, Francisco José Heredia<sup>†</sup>, Gerard  
Downey<sup>‡,\*</sup>

<sup>†</sup> Food Colour & Quality Laboratory, Department of Nutrition & Food Science,  
Universidad de Sevilla, Facultad de Farmacia, 41012 Sevilla, Spain.

<sup>‡</sup> Teagasc, Ashtown Food Research Centre, Dublin 15, Dublin, Ireland.

<sup>§</sup> Horticulture Development Unit, Teagasc Research Centre, Kinsealy, Dublin 17,  
Ireland.

**\* Corresponding author:** Gerard Downey

**Phone:** (+353 1) 805 9500

**Fax:** (+353 1) 805 9550

**E-mail:** Gerard.Downey@teagasc.ie

1    **ABSTRACT**

2    The potential of visible-near infrared spectroscopy to determine selected individual and  
3    total glucosinolates in broccoli has been evaluated. Modified partial least squares  
4    (MPLS) regression was used to develop quantitative models in order to predict  
5    glucosinolate contents. Both the whole spectrum and different spectral regions were  
6    separately evaluated to develop the quantitative models, in all cases the best results were  
7    obtained using the NIR zone between 2000 and 2498 nm. These models have been  
8    externally validated for the screening of glucoraphanin, glucobrassicin, 4-  
9    methoxyglucobrassicin, neoglucobrassicin and total glucosinolates contents. In addition,  
10   discriminant partial least squares was used to distinguish between two possible broccoli  
11   cultivars and showed a high degree of accuracy. In the case of the qualitative analysis,  
12   best results were obtained using the whole spectrum (i.e. 400-2498 nm) with a correct  
13   classification rate of 100% in external validation being obtained.

14

15   **KEYWORDS:** Glucosinolates; broccoli; visible spectroscopy; near infrared  
16   spectroscopy; chemometrics

17

## 18 INTRODUCTION

19 Plant bioactive compounds, commonly referred to as phytochemicals, are components  
20 that can affect cells in ways other than by provision of nutrients. Some of them may  
21 have functional properties, meaning that they provide health benefits beyond basic  
22 nutrition. It has been suggested that the health-promoting properties of plant foods are  
23 due to their content of bioactive components with health-promoting effects<sup>1-3</sup>.

24 Broccoli (*Brassica oleracea* L. var *Italica*) has been produced and consumed in Europe  
25 since early times and is a characteristic constituent of European diets. This vegetable  
26 contains significant amounts of glucosinolates, which are small organic molecules not  
27 essential in the primary metabolism of the plant but with specific functions within its  
28 secondary metabolism. Although their role in plants is unclear, their potent odour and  
29 pronounced taste suggests a role in herbivore and microbial defence. Structurally,  
30 glucosinolates ( $\beta$ -thioglucoside-N-hydroxysulfates) (**Figure 1A**) are characterised by  
31 the presence of nitrogen and sulphur groups and they are derived from glucose and an  
32 amino acid<sup>1</sup>. Glucosinolates are not bioactive until they have been enzymatically  
33 hydrolysed to the associated isothiocyanate by an endogenous myrosinase enzyme that  
34 may be released by disruption of the plant cell through harvesting, processing or  
35 mastication. Epidemiological studies have consistently reported a reduction in incidence  
36 of chronic diseases such as cancer and myocardial infarction as a result of induction of  
37 detoxifying enzymes and reduction of oxidative stress, although the anti-nutritive  
38 effects of both glucosinolates and hydrolysis products have also been studied<sup>1,4,5</sup>. It is  
39 therefore important to characterise the content of bioactive compounds in new or re-  
40 introduced cultivars.

41 To this end, the use of rapid methods of analysis with minimal or no sample preparation  
42 could be important. Such a capability would significantly enhance the ability of primary

43 processors to select samples for the market based on a high content of bioactive  
44 compounds in a rapid and inexpensive manner.

45 Near infrared spectroscopy (NIRS) provides fast and non-destructive analysis. It allows  
46 qualitative and quantitative analysis to be performed in different matrices thereby  
47 reducing costs when compared to wet chemical analysis and without generating waste.  
48 In the recent past, the use of NIR spectroscopy in the determination of bioactive  
49 compounds in foods and plant natural products has increased considerably <sup>6, 7</sup>. With  
50 particular regard to glucosinolates, it has been used to determine these compounds in  
51 both seeds <sup>8-13</sup> and leaves <sup>14, 15</sup> from a wide range of *Brassica* species. The standard  
52 errors of the glucosinolate's predictions reported in the aforementioned studies ranged  
53 from 2.65 to 15.65  $\mu\text{mol g}^{-1}$  DM for three cultivars of rapeseeds <sup>13</sup> and Indian mustard  
54 seeds <sup>11</sup> respectively. The results reported in all these works reveal that NIRS is able to  
55 predict glucosinolates in a wide range of matrices. Therefore, it may be expected that  
56 this technique coupled with chemometric tools could provide an alternative method to  
57 undertake the analysis of glucosinolates in broccoli.

58 The aim of this study was to evaluate the potential of visible-NIR technology to  
59 determine both selected individual (glucoraphanin, glucobrassicin, 4-  
60 methoxyglucobrassicin, neoglucobrassicin) and total glucosinolates in broccoli. In  
61 addition to this, its potential for the discrimination between two broccoli cultivars was  
62 evaluated. To our knowledge, this is the first time that vis-NIRS technology has been  
63 applied to broccoli for these purposes. The work is part of a wider study aimed at  
64 maximising the concentrations of phytochemicals in vegetables by adopting a farm to  
65 fork approach. The projected final output of the wider study is the production of  
66 vegetables with optimal levels of selected phytochemical groups.

## 67 MATERIAL AND METHODS

## 68 **Samples**

69 Samples were generated at Teagasc, Kinsealy Research Centre, located in the north of  
70 County Dublin, Ireland (53° 25' 10" N Lat 6° 10' 45" W). The soil is a grey brown  
71 podzolic and presents a loam to clay loam topsoil. The experimental design was 2<sup>3</sup>  
72 factorial split plot design with four plot replicates. Briefly, eight treatments in 5.5m x  
73 3.4 m plots were applied with four replicates combining conventional or organic soil  
74 management with organic or conventional pest control. The four combinations were  
75 sown with Belstar or Fiesta broccoli cultivars. **Table 1** shows the different agricultural  
76 managements applied in this study. The aforementioned factorial trial was carried out  
77 during the 2009 and 2010 years and provided 64 samples.

78 Broccoli samples were harvested, frozen and freeze-dried in a large scale freeze drier  
79 (Frozen in Time Ltd. York, UK). The specimen chamber was kept below 0 °C during  
80 the whole freeze-drying process. Once freeze-dried, samples were milled (Blixer 4,  
81 Robotcoupe, France), vacuum packed in polypropylene bags and stored at -20 °C prior  
82 to analysis. Two aliquots were taken from each sample, one for the micellar  
83 electrokinetic capillary chromatography (MEKC) analysis and the other for visible-near  
84 infrared analysis.

## 85 **Chemical Analysis**

86 *Sulphatase Purification Procedure.* Sulphatase (Type H-1 from *Helix pomatia*, Sigma,  
87 MO, USA) was purified by dissolving the sulphatase powder (70 mg) in deionised  
88 water (3 mL) and adding ethanol (3 mL). This solution was centrifuged (12000 rpm, 10  
89 min, room temperature) and ethanol (9 mL) was added to the supernatant after which  
90 the solution was centrifuged again (12000 rpm, 10 min, room temperature). The  
91 resulting pellet was dissolved in deionised water (2 mL) and this sulphatase solution  
92 was subsequently passed through a 0.5 mL DEAE Sephadex A-25 and a 0.5 mL SP

93 Sephadex C-25 column. This solution was collected in a vial and kept at -20 °C until  
94 use.

95 *Sampling and Extraction.* Freeze-dried broccoli powder was extracted using pressurised  
96 liquid extraction with an ASE200 instrument (Dionex; Sunnyvale, CA, USA) with  
97 attached solvent controller. Extraction of 1.00 g freeze-dried broccoli was carried out in  
98 22 mL steel cartridges. Extraction conditions were slightly modified from Mohn *et al*<sup>16</sup>.  
99 In brief, preheat time: 1 min; static extraction per cycle: 5 min; flush: 60% of cell  
100 volume; purge: 60 s with nitrogen; pressure: 120 bar. Glucotropaeolin (Phytolab GmbH,  
101 Vestenbergsgreuth, Germany) was used as an internal standard - 100 µL of a solution of  
102 0.5 µmol of potassium salt of glucotropaeolin in Milli-Q water was spiked onto the  
103 sample packed in the ASE cell. The collected extract (ca 30 mL) was dried under a  
104 constant N<sub>2</sub> flow and re-dissolved in deionised water (7 mL). An aliquot (1 mL) of this  
105 crude extract was applied to a DEAE Sephadex A-25 column (0.5 mL) and the unbound  
106 material was removed by washing with deionised water (2 x 1 mL) and sodium acetate  
107 buffer (2 x 0.5 mL; 20 mM, pH 5.0). After washing, the purified sulphatase (75 µL) was  
108 added and the columns were incubated overnight at room temperature. After overnight  
109 incubation, the desulphoglucosinolates (dGLS) were eluted from the columns with  
110 deionised water (3 x 1 mL). The collected eluate was dried under constant N<sub>2</sub> flow, re-  
111 dissolved in deionised water (200 µL) and centrifuged prior to analysis. An aliquot of  
112 50 µL was used for the MEKC analysis.

113 *Micellar Electrokinetic Capillary Chromatography.* Analyses were performed using a  
114 CE capillary electrophoresis system (Agilent, Waldbronn, Germany) equipped with  
115 diode array detector. All separations were performed on a fused silica capillary (Agilent,  
116 Stevens Creek, CA; 75 µm ID, 64.5 cm total length, 56 cm effective length). Samples  
117 were injected from the anodic end of the capillary (vacuum injection, 50 mbar, 1 s). The

118 separation buffer consisted of sodium chlorate (250 mM) and boric acid (200 mM) at  
119 pH 8.5; the separation was carried out at 12 kV and 60 °C. The capillary was  
120 conditioned between each run sequentially with 1.0 M NaOH (3 min), 0.1 M NaOH (1  
121 min), water (1 min) and separation buffer (5 min). Detection was performed on column  
122 at 230 and 280 nm. Data processing was carried out with 3D-CE Chemstation software  
123 (Agilent Waldbronn, Germany). The quantity of the dGLS was estimated as the average  
124 of quantities calculated from the internal standards, taking into account the relative  
125 response factors revised by Clarke *et al*<sup>17</sup>.

### 126 **Near Infrared Spectroscopy Analysis**

127 Each aliquot of freeze-dried broccoli powder was tempered at room temperature for 16  
128 hours prior to spectroscopy analysis. A Foss NIRSystems 6500 spectrometer (NIR  
129 Systems Inc., Silver Spring MD) was used; samples were placed in small ring cups (3  
130 cm diameter screen size) with a disposable backing disc and spectral measurements  
131 were made in reflectance mode over the 400-2498 nm wavelength range. Spectra were  
132 recorded in duplicate at intervals of 2 nm with 10 scans being performed for the  
133 reference tile and 25 for samples; the sample cell was rotated through 180 degrees  
134 between spectral collections. To minimise sampling error, all the samples were analysed  
135 in duplicate with re-filling of the ring cup. Therefore, the four recorded spectra of each  
136 sample were averaged to obtain each sample spectrum. Samples were scanned in  
137 random order. The spectrum of sinigrin standard powder ((-)-sinigrin hydrate from  
138 horseradish, Sigma-Aldrich Ireland Ltd, Arklow, Ireland) was also recorded in  
139 reflectance mode over the wavelength range 1100-2498 nm at intervals of 2 nm. The  
140 software used was Win ISI<sup>®</sup> (v1.50) (Infrasoft International, LLC, Port. Matilda, PA,  
141 USA). This software allowed not only the instrument control and spectral acquisition  
142 but also the data pre-treatment and development of qualitative and quantitative models.



143 From the whole data set, 25% of the samples (16 samples) were randomly allocated into  
144 the validation set and the remaining 75% (48 samples) into the calibration set.

#### 145 **Chemometric Techniques**

146 A supervised pattern recognition technique-Discriminant Partial Least Squares (DPLS)-  
147 was used for qualitative analysis. The calibration was conducted by performing a  
148 regression on the spectral information and the associated dummy sample group values,  
149 in this case defined as 1 (variety 1) or 2 (variety 2). The regression method applied to  
150 this procedure was MPLS, which is a modification of a normal PLS 1<sup>18,19</sup>.

151 Prior to quantitative analysis, an unsupervised pattern recognition technique, principal  
152 component analysis (PCA), was used in order to provide information about the latent  
153 structure of spectral data. This method provides not only information related to spectral  
154 outliers and the distribution of samples in the newly-created space but is also an  
155 important source of knowledge with which to create cross-validation groups used in the  
156 calibration process. PCA is also a useful tool to identify whether unknown samples  
157 belong to the spectral space created by the samples from which the equations were  
158 developed. Should this not be the case, the equations should not be used to make any  
159 prediction<sup>20,21</sup>. Using the raw spectral data and testing different spectral pre-treatments  
160 and allocating the corresponding MEKC-DAD glucosinolate values to each sample,  
161 calibrations were performed by modified partial least squares regression (MPLS). In this  
162 method, the set of calibration samples is divided into a series of subsets in order to  
163 perform cross-validation to set the number of PLS factors, reduce the likelihood of  
164 overfitting<sup>20</sup> and remove chemical outliers. Using the  $T \geq 2.5$  criterion, samples that  
165 presented a high residual value when they were predicted were eliminated from the set.  
166 Finally, validation errors are combined into a single figure, the standard error of cross-  
167 validation (SECV).

168 Spectral pre-treatments are usually applied to NIR raw data; scattering effects were  
169 removed using multiplicative scatter correction (MSC), standard normal variate (SNV),  
170 and detrending<sup>22, 23</sup>. Moreover, the effect of derivatisation and variations in spectral  
171 ranges were tested in the development of the NIRS calibrations.

## 172 **RESULTS AND DISCUSSION**

### 173 **Chemical Analysis**

174 Glucosinolates were determined by MEKC-DAD. The common glucosinolate skeleton  
175 structure is shown in **Figure 1A**, leading to the expectation that their spectral signatures  
176 should also be very similar. Sinigrin's reflectance spectrum (**Figure 1B**) shows  
177 considerable spectral detail in the wavelength range above about 2000 nm; it was the  
178 only pure glucosinolate available for spectral analysis but, given that its structure is  
179 similar to the others being analysed in these broccoli samples, it is expected that all  
180 glucosinolates will have similar information above 2000 nm. **Table 2** shows the range,  
181 mean value and standard deviation of the selected individual glucosinolates and the total  
182 glucosinolates in the broccoli samples analysed. These values were used in the  
183 development of the quantitative models as reference values. It is noticeable that the  
184 precision of the laboratory methods is quite poor with standard deviation between  
185 replicate figures of between 15.9% (neoglucobrassicin) and 27.2% (glucoraphanin) of  
186 mean contents.

### 187 **Near Infrared Spectroscopy Analysis and Chemometric Techniques**

188 *Near Infrared Spectra.* **Figure 1** shows the average and standard deviation spectra of  
189 broccoli powders over the 400-2498 nm range (**Figure 1C**) together with the sinigrin  
190 powder spectrum between 1100-2498 nm (**Figure 1B**).

191 Standard deviation spectra have been multiplied by a factor of 10 for display reasons.  
192 Spectral intensities were low and well within the linear response range of the instrument  
193 detector range; little noise was evident in these traces. A strong feature of the sample  
194 spectra was the absorbance pattern in the visible wavelength range i.e. 400-780 nm.  
195 This arose from the pigment remaining in the freeze-dried samples, all of which  
196 displayed a green tinge.

197 **Figure 1D** shows the second derivative spectra of broccoli and sinigrin and presents  
198 considerable spectral similarities over the 2000-2498 nm wavelength range. It may be  
199 expected that this range would be useful for quantitative purposes of glucosinolates.

200 *Qualitative analysis.* An SNV 2,4,4,1 spectral pre-treatment was applied to whole vis-  
201 NIR spectra of samples from the calibration set and then a principal component analysis  
202 was carried out. **Figure 2** shows the scores of the broccoli samples in the space defined  
203 by the first and second principal components which described 52.3% (PC1) and 23.7%  
204 (PC2) of the variability in the data. In this plot, slight differences between the 2009 and  
205 2010 harvest years (**Figure 2A**) and also between the Belstar and Fiesta cultivars  
206 (**Figure 2B**) are apparent. The main difference observed in these plots was between  
207 harvest years although the separation between 2009 and 2010 samples was not  
208 complete. This trend in harvest year locations was discernible on the basis of PC2.  
209 Cultivars were completely overlapped in this hyperspace.

210 D-PLS modelling of the groups was carried out initially using the complete vis-NIR raw  
211 spectra and one dummy variable, the values of which were defined as 1 (variety  
212 1=Belstar) or 2 (variety 2=Fiesta). The model was developed using 6 PLS factors and  
213 presented an RSQ of 0.752, SEC of 0.27 and SECV 0.34. The spectral regions between  
214 600-700 nm and 2000-2498 nm showed important contributions to the model loadings  
215 (**Figure 3A**) and are mainly related to green pigments such as chlorophyll<sup>14, 15, 24</sup> and C-

216 H, O-H and N-H overtones present in a number of bonds<sup>25, 26</sup>. The DPLS developed  
217 model predicted a dummy value for each sample and then the samples were allocated  
218 according to their predicted values  $\pm 0.5$  in the corresponding cultivar with a dummy  
219 variable breakpoint of 1.5. The predicted dummy variable plot for the qualitative model  
220 is shown in **Figure 3B**. Samples of Belstar cultivar cluster around a dummy value of  
221 1.0, while those belonging to Fiesta cultivar around 2.0. A prediction rate of 100%  
222 samples correctly classified was obtained in external validation. This demonstrates that  
223 vis-NIR spectroscopy coupled to discriminant DPLS analysis permit a clear  
224 differentiation between the two aforementioned broccoli cultivars. Regarding the  
225 studied compound, the classification of samples according to their cultivar could be  
226 useful since they present significant differences in the 4-methoxyglucobrassicin  
227 ( $p < 0.001$ ) and neoglucobrassicin ( $p < 0.001$ ) contents.

228 *Quantitative analysis.* Although different spectral ranges and the whole spectrum have  
229 been separately evaluated, in all cases the best results were obtained using the  
230 wavelength range 2000-2498 nm. As part of the quantitative analysis, a SNV (2,4,4,1)  
231 spectral pre-treatment was applied to this spectral range of samples in the calibration set  
232 and then principal component analysis was carried out in order to look for spectral  
233 outliers and create cross-validation groups. Overall, the spectral variability explained  
234 was 98% using 13 principal components and Mahalanobis distances for each sample  
235 were calculated. Samples were ranked in order of their H (Mahalanobis) distance from  
236 the mean spectrum of the entire sample set and the  $H > 3$  criterion was applied. No H-  
237 outliers were found.

238 The average spectrum of the best of the different mathematically pre-treated spectra are  
239 shown in **Figure 4**. The statistical parameters of the final calibration equations are  
240 shown in **Table 3** where N is the number of samples used to obtain the calibration

241 equation after eliminating samples for chemical reasons (T criterion). The best of the  
242 different mathematical treatments, concentration range, and standard deviations are also  
243 shown. We checked the robustness of the method by applying NIRS technology to 16  
244 samples that did not belong to the calibration group. **Table 3** also shows the results  
245 obtained in the external validation and the SEP values are presented. The percentage of  
246 error for total glucosinolates is similar to the errors previously reported for other  
247 matrices such as Indian mustard seeds<sup>11</sup>.

248 The magnitude of these errors indicate the potential of the NIR spectroscopy for  
249 prediction of glucosinolates in freeze-dried broccoli and is satisfactory for the screening  
250 of glucoraphanin, glucobrassicin, 4-methoxyglucobrassicin, neoglucobrassicin and total  
251 glucosinolates. However, individual glucosinolates biosynthesis may follow essentially  
252 the same course so inter-correlations among them could be expected. It is not possible  
253 to ascertain if the results of NIRS models for predicting the composition of these  
254 glucosinolates were due to their real absorbance or the correlation between them.

255 **Figure 5** shows the loading and the beta regression coefficients plots of the MPLS  
256 model for total glucosinolates. The spectral regions around 2072 nm, 2220 nm, 2300 nm  
257 and 2446 nm show important contributions to the model loadings. These could be  
258 related to N-H bonds (2072 nm), to C-H bonds (2220 nm), to first overtones of O-H  
259 alcohol functional group, to stretching-bending of CH-CH<sub>2</sub> bonds and C-O bonds (2300  
260 nm) and to C-N bonds (2446 nm). These can be attributed to the chemical structure of  
261 the compounds analyzed<sup>25, 26</sup>. This confirms previous studies that showed important  
262 contributions in the aforementioned spectral zones for determining glucosinolates in  
263 other matrices<sup>11, 14, 15, 27</sup>.

264 The potential of vis-NIRS for determination of glucosinolates in freeze-dried broccoli  
265 was examined. The procedure reported here presents a good potential for a fast and

266 reasonably inexpensive screening of almost all individual glucosinolates and total  
267 glucosinolates present in broccoli and also to distinguish between broccoli cultivars.  
268 Although different spectral ranges and the whole spectrum have been separately  
269 evaluated to develop the quantitative models, in all cases the best results (i.e. the highest  
270 RSQ and the lowest SECV and SEP values) were obtained using the NIR region  
271 between 2000 and 2498 nm. In the case of the qualitative analysis, best results were  
272 obtained using the whole spectrum (i.e. 400-2498 nm). Nonetheless, a comprehensive  
273 study should be made in order to evaluate other factors such as different production  
274 areas and varieties, in the complete development of these models.

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281

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361



**Figure captions**

**Figure 1.** (A) Glucosinolate skeleton. (B) Sinigrin powder spectrum (1100-2498 nm). (C) Average and standard deviation spectra (400-2498 nm) of broccoli powders. (D) 2<sup>nd</sup> derivative spectra of the broccoli and sinigrin (1100-2000 nm).

**Figure 2.** Score plot of broccoli samples (calibration set) in the space defined by PC1 and PC2.

**Figure 3.** (A) Loading plot of the DPLS model. (B) Dummy variable plot of broccoli samples in the external validation of the DPLS method.

**Figure 4.** Average spectrum of different mathematical pre-treatments applied to broccoli powder spectra (2000-2498 nm).

**Figure 5.** (A) Loading and (B) beta regression coefficients plots of the MPLS model for total glucosinolate prediction.

**Table 1.** Different Agricultural Managements Applied in this Study.

	Organic	Conventional
Soil treatment	<ul style="list-style-type: none"> <li>• Four year rotation ley: red clover-broccoli-onion-carrot</li> <li>• Additional organic fertilisation as indicated by soil test</li> <li>• Winter cover crop</li> </ul>	<ul style="list-style-type: none"> <li>• No rotation, plots, randomly allocated each year</li> <li>• Mineral fertilisers as indicated by soil test</li> <li>• No ley crop</li> <li>• No winter crop</li> </ul>
Pest control	<ul style="list-style-type: none"> <li>• Certified organic seed</li> <li>• Refuge area</li> <li>• <i>Brassica</i> collars on broccoli</li> <li>• Certified organic chemicals (eg. garlic spray)</li> <li>• Weed control by mechanical methods</li> </ul>	<ul style="list-style-type: none"> <li>• Chemically treated seed</li> <li>• Chemical weed control–herbicides</li> <li>• Chemical pest control–fungicides and insecticides</li> </ul>

**Table 2.** Statistical Descriptors of Glucosinolate Contents Determined by MEKC-DAD ( $\mu\text{mol g}^{-1}$  DM).

	GBS <sup>a</sup>	GRA <sup>b</sup>	4MGBS <sup>c</sup>	NGBS <sup>d</sup>	TGS <sup>e</sup>
Min (n=64)	1.60	1.09	0.09	0.74	4.74
Max (n=64)	6.83	4.82	0.55	4.45	15.10
Mean (n=64)	3.92	2.83	0.28	1.97	9.53
SD (n=64)	1.09	0.87	0.10	0.86	1.87
SD replicates	0.88	1.01	0.08	0.59	1.80

<sup>a</sup>GBS: glucobrassicin; <sup>b</sup>GRA: glucoraphanin; <sup>c</sup>4MGBS: 4-methoxyglucobrassicin; <sup>d</sup>NGBS: neoglucobrassicin; <sup>e</sup>TGS: total glucosinolates

**Table 3.** Calibration Statistical Descriptors for the Models Developed in the NIR Zone Close to 2000-2498 nm.

Spectral pre-treatments	Compounds	T outliers	PLS factors	N <sup>a</sup>	(μmol g <sup>-1</sup> DM)					(μmol g <sup>-1</sup> DM)		
					Mean	Est. Min	SD <sup>b</sup>	Est. Max	SEC <sup>c</sup>	RSQ <sup>d</sup>	SECV <sup>e</sup>	SEP <sup>f</sup>
Detrend 1,4,4,1	GBS <sup>g</sup>	2	7	46	3.93	0.68	1.09	7.18	0.35	0.89	0.52	0.74
Detrend 1,4,4,1	GRA <sup>h</sup>	1	2	47	2.86	0.26	0.87	5.46	0.67	0.40	0.71	0.75
SNV 1,4,4,1	4MGBS <sup>i</sup>	0	5	48	0.28	0.00	0.11	0.62	0.06	0.69	0.07	0.05
MSC 2,8,6,1	NGBS <sup>j</sup>	0	5	48	1.93	0.00	0.86	4.51	0.48	0.68	0.66	0.74
None 2,10,10,1	TGS <sup>k</sup>	3	5	45	9.40	4.04	1.79	14.76	0.93	0.73	1.11	1.44

<sup>a</sup>N: number of samples (calibration set); <sup>b</sup>SD: standard deviation; <sup>c</sup>SEC: standard error of calibration; <sup>d</sup>RSQ: coefficient of determination;

<sup>e</sup>SECV: standard error of cross-validation; <sup>f</sup>SEP: standard error of prediction; <sup>g</sup>GBS: glucobrassicin; <sup>h</sup>GRA: glucoraphanin;

<sup>i</sup>4MGBS: 4-methoxyglucobrassicin; <sup>j</sup>NGBS: neoglucobrassicin; <sup>k</sup>TGS: total glucosinolates

Figure 1.

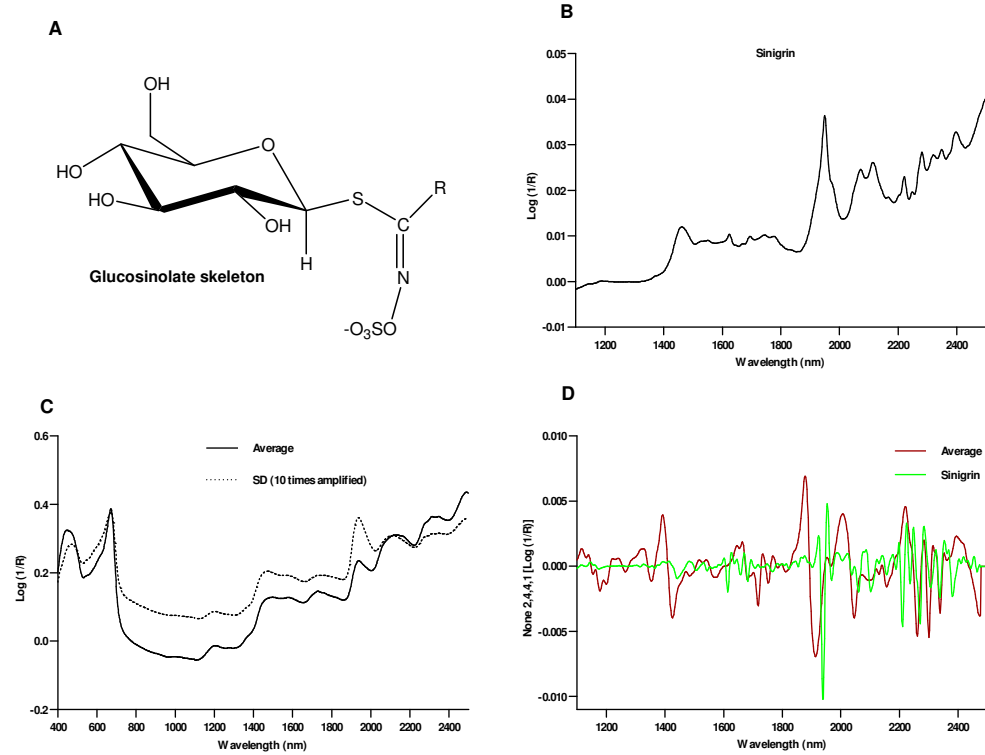


Figure 2.

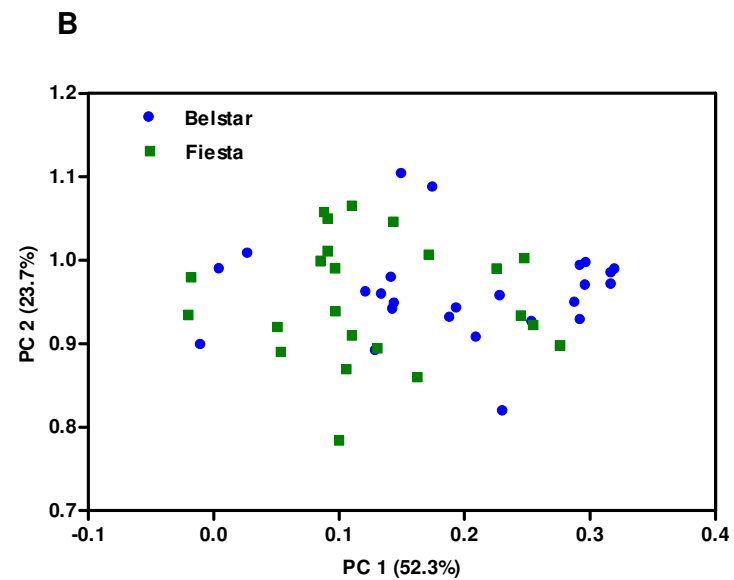
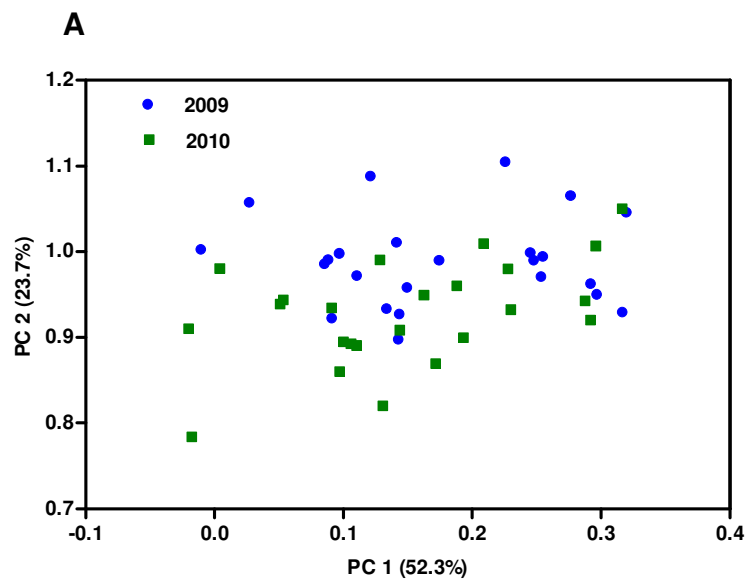


Figure 3.

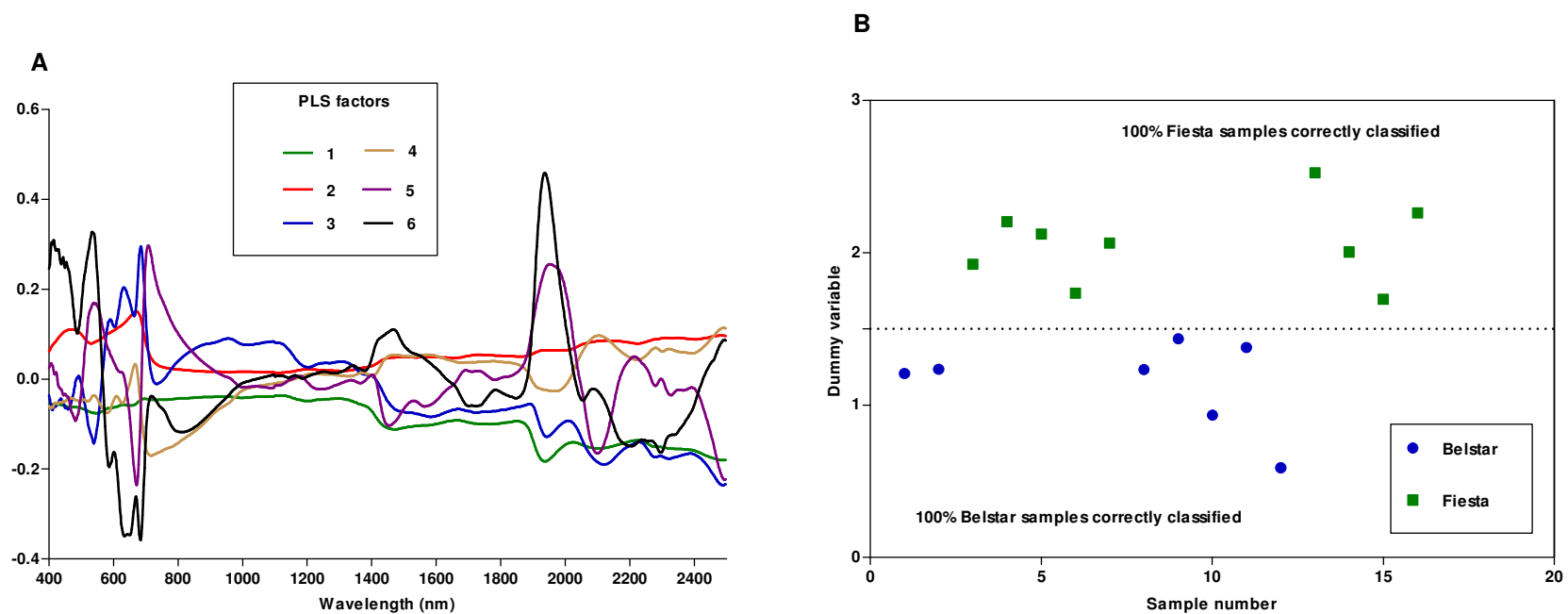


Figure 4.

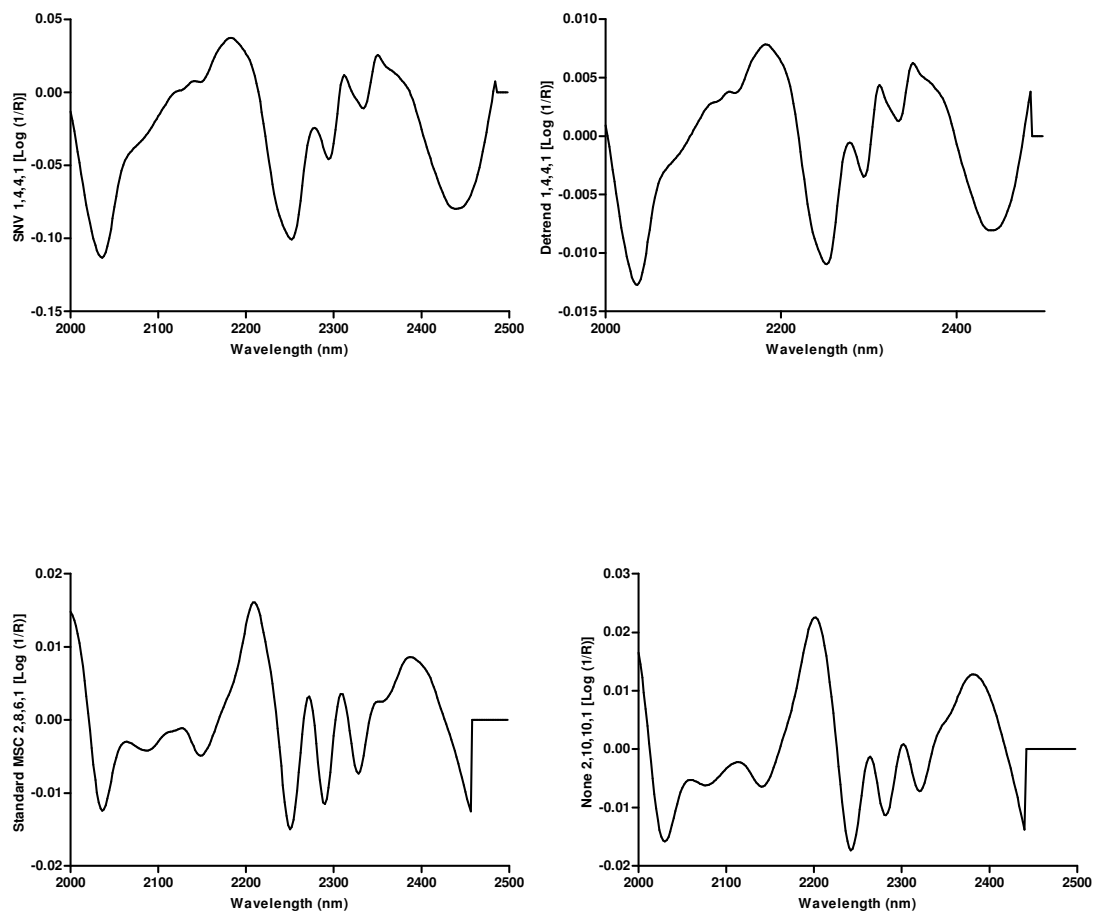
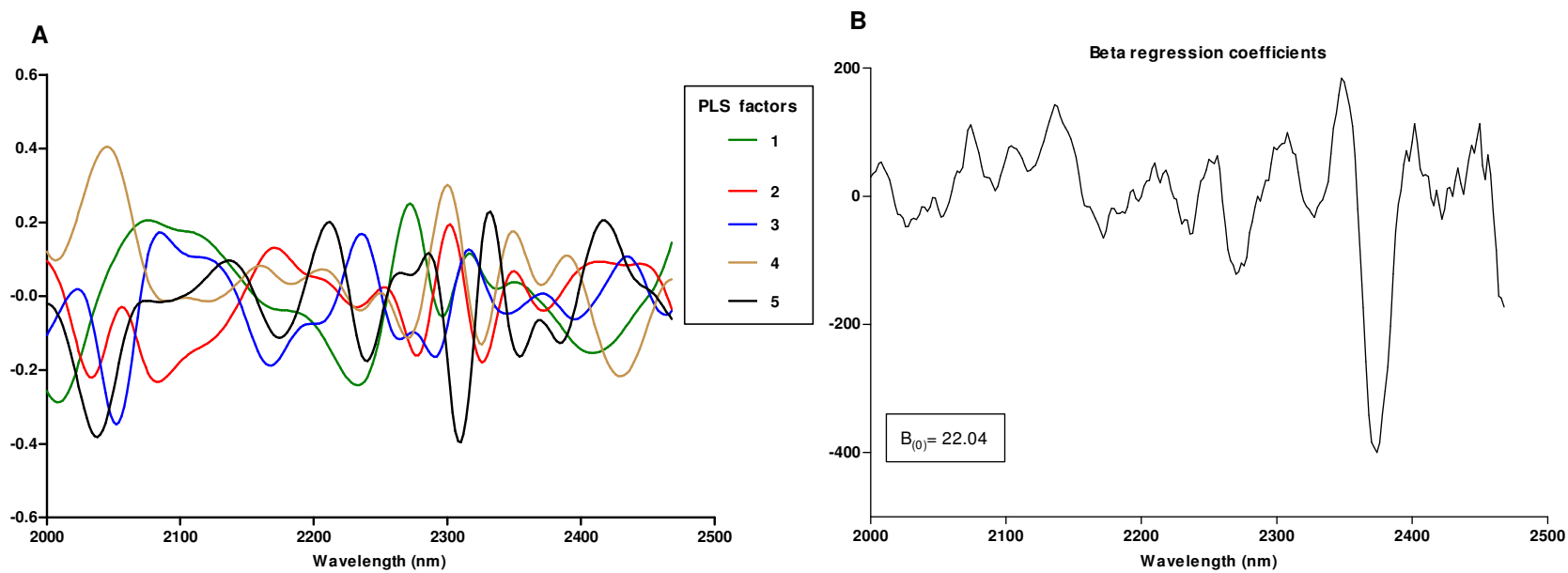




Figure 5.



## Graphic for table of contents

