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Quantification of leghaemoglobin content in pea nodules based on near infrared hyperspectral imaging spectroscopy and chemometrics

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Leghaemoglobin content in nodules is closely related to the amount of nitrogen fixed by the legume-rhizobium symbiosis. It is, therefore, commonly measured in order to assess the effect of growth-promoting parameters such as fertilisation on the symbiotic nitrogen fixation efficiency of legumes. The cyanmethaemoglobin method is a reference method in leghaemoglobin content quantification, but this method is time-consuming, requires accurate and careful technical operations and uses cyanide, a toxic reagent. As a quicker, simpler and non-destructive alternative, a method based on near infrared (NIR) hyperspectral imaging was tested to quantify leghaemoglobin in dried nodules. Two approaches were evaluated: (i) the partial least squares (PLS) approach was applied to the full spectrum acquired with the hyperspectral device and (ii) the potential of multispectral imaging was also tested through the preselection of the most relevant wavelengths and the building of a multiple linear regression model. The PLS approach was tested on mean spectra acquired from samples containing several nodules and acquired separately from individual nodules. Peas (*Pisum sativum* L.) were cultivated in a greenhouse. The nodules were harvested on four different dates in order to obtain variations in leghaemoglobin content. The leghaemoglobin content measured with the cyanmethaemoglobin method in fresh nodules ranged between 1.4 and 4.2 mgleghaemoglobin gr⁻¹ fresh nodule. A PLS regression model was calibrated on leghaemoglobin content measured with the reference method and mean NIR spectra of dried nodules acquired with a hyperspectral imaging device. On a validation dataset, the PLS model predicted the leghaemoglobin content in nodule samples well (*R*² = 0.90, root mean square error of prediction = 0.26). The multispectral approach showed similar performance. Applied to individual nodules, the PLS model highlighted a wide variability of leghaemoglobin content in nodules harvested from the same plant. These results show that NIR hype

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Introduction

Legume plants can fix atmospheric nitrogen thanks to a symbiosis with bacteria. The leghaemoglobin protein is synthesised by the host plant within the cytoplasm of cells located in the nitrogen-fixing zone of nodules, the special organ of legumes that contains the nitrogen-fixing bacteria. Leghaemoglobin has a high affinity with oxygen and keeps the $\rm O_2$ concentration in an optimal range to protect nitrogenase enzyme against oxygen damage and to supply bacterial respiration. The nitrogenase enzyme is contained in the nodule and transforms $\rm N_2$ into NH₃, a form of nitrogen which is metabolically available for plants. In the case of pea (*Pisum sativum L.*), the bacteria involved in this transformation is *Rhizobium leguminarosum*.

Leghaemoglobin in root nodules is commonly quantified in order to assess the effects of plant growth-promoting parameters [such as fertilisation or ultraviolet (UV) exclusion of solar radiation] on symbiotic nitrogen fixation efficiency. $^{1.6.7}$ The concentration of leghaemoglobin is affected by several factors, such as the age of the nodules and the content of nutrients like S or N in the growing substrate or plant health. $^{1.6.8}$ Its concentration in root nodules is closely correlated with the amount of nitrogen fixed in the association between the plant and the bacteria. $^{5.9}$ Low leghaemoglobin content in nodules leads to low $\rm N_2$ fixation due to higher $\rm O_2$ concentration and inhibition of the nitrogenase enzyme. $^{1.4}$

Common methods of leghaemoglobin quantification require prior extraction of leghaemoglobin from nodules before any possible quantification of its concentration. This extraction step is time-consuming, destructive and sensitive to operator ability and needs several chemical reagents. Moreover, several nodules are needed and leghaemoglobin quantification is, to the best of our knowledge, not possible in an individual nodule. Among the methods used to quantify leghaemoglobin, the most popular one is based on the cyanmethaemoglobin method, also used in medicine to quantify haemoglobin in blood. As leghaemoglobin is structurally and chemically similar to human haemoglobin, the method was transposed to legume haemoglobin as

described by Wilson and Reisenaeur. ⁹ The principle of this method is the conversion of haemoglobin into cyanmethaemoglobin by the addition of potassium cyanide and ferricyanide, whose absorbance is measured at 540 nm in a spectrophotometer against a standard solution. 10 This method is considered as a reference method because it is accurate, cheap, sensitive to small variations in haemoglobin concentration, reliable, reproducible and requires a minimum of specialised equipment and reagents. 9-12 Nevertheless, the multiple operations required in this method make it time-consuming and thus unsuitable for large-scale field studies. Furthermore, it requires accurate and careful technical operations, which increase the risk of imprecision and errors. 10 Finally, this method uses cyanide, a toxic reagent. 11

In order to avoid the limitations of the cyanmethaemoglobin method in leghaemoglobin extraction and content measurement, the use of near infrared (NIR) hyperspectral imaging combined with chemometrics was studied as a new way to quantify leghaemoglobin in nodules. NIR hyperspectral imaging is the combination of NIR spectroscopy and imaging technologies. This technology provides spectral and spatial information simultaneously, i.e. thousands of spectra can be obtained for each sample and give a complete picture of the chemical compounds' distribution at the pixel level. This technology requires the use of chemometrics, a chemical discipline using mathematics and statistics, in order to extract relevant information from the measurement procedures. ^{13,14}

The aim of the present study was to predict the leghaemoglobin concentration of nodules on the basis of their NIR spectra. Two approaches were evaluated. First, the partial least squares (PLS) approach was applied to the full spectrum acquired with the hyperspectral device. Second, the potential of multispectral analysis was also tested through the preselection of the most responsive wavelengths and the application of a multiple linear regression model. Both procedures were tested on a standard calibration–validation approach.

Materials and method

Nodule production

To produce nodules that covered a large range of leghaemoglobin concentrations, a dedicated protocol was established taking into account that leghaemoglobin content in pea is affected by nitrogen fertilisation and varies over time with plant growth. Two types of fertilisation (no fertilisation and nitrogen fertilisation) were applied at seeding time of pea plants. Under both fertilisation treatments, nodules were harvested at four different dates during plant growth. Each one of the eight treatments (fertilisation × harvest date) was replicated four times.

Nodules were produced in pots (20 cm in diameter, 15 cm high) in the greenhouses of Gembloux Agro-Bio Tech (University of Liège, Belgium). Five pea seeds were sown in each pot containing field soil and sand (5 kg of substrate per pot with a mass ratio of ¾ field soil and ¼ sand). The field soil was collected in a field (loam type) where peas had been cultivated three months earlier to ensure the presence of rhizobium in the substrate. Seeds of the Dove variety were used. They were protected with WAKIL fungicide (10% Cymoxanil, 5% Fludioxonil and 17% Metalaxyl-m). The fertiliser was applied in the top layer of pot soil at seeding time. Nitrate fertilisation consisted of one input of 0.49 g ammonitrate per pot (equivalent to 80 kg N ha⁻¹). Plants were cultivated between 12 October and 30 November under a 24h photoperiod (daylight and artificial light complement). Being exposed to aphids 35 days after seedling emergence, the plants were protected with Pirimor (50% Pirimicarbe). The temperature was measured in the vicinity of the plants throughout their development. After germination, three plants were kept in each pot.

The first nodule harvest occurred 25 days after sowing (on 6 November). At this time, the plants had accumulated 500 heat units (basal temperature = 0 $^{\circ}$ C) and had produced seven leaves. The second harvest occurred when the plants had accumulated 630 heat units, the third after 760 heat units and the fourth after 820 heat units, 48 days after sowing. Approximately one week separated each harvest.

At each harvest date, four pots were randomly selected in each fertilisation modality. In order to quantify plant development, the number of nodules present on each plant root system and the aerial dry biomass were measured. Roots were manually extracted from soil with tap water and nodules were separated from roots with tweezers. For each pot, a sample of 30 nodules was

dried at 60 °C and then kept in ambient conditions until image acquisition. Other nodules were saved in falcon tubes, dipped in liquid nitrogen and kept at -80 °C. When the number of nodules harvested on a plant was below 60, only 10 nodules were dried in order to save enough nodules for the chemical quantification.

Leghaemoglobin quantification

Leghaemoglobin content in frozen nodules was measured the day after each harvest. The assays were based on the cyanmethaemoglobin method described in detail by Wilson and Reisenauer. ⁹ This method is based on spectrophotometric measurement and uses Drabkin's solution.¹⁷ Briefly, Drabkin's solution was prepared with 52 mg KCN, 198 mg K₈Fe(CN)₆ and 1 g NaHCO₃ dissolved in water and made up to 1000 mL. To extract leghaemoglobin, frozen nodules were crushed in liquid nitrogen and 100 mg of crushed nodules was transferred to a 2 mL centrifuge tube with 0.6 mL Drabkin's solution. The mixture was centrifuged for 15 min at $500 \times g$ and $4 \,^{\circ}$ C. The supernatant was transferred to a 2mL flask. The solid phase was extracted and centrifuged twice more. Supernatants were combined, made up to 2 mL with Drabkin's solution and centrifuged for 30 min at 20,000 × g and 4 °C. Absorbance of the cleared supernatant was read against Drabkin's solution in a 1.5 mL cell at 540 nm with a UV-visible spectrophotometer (UV-1650 PC, Shimadzu, Japan).

A reference curve was computed each day of measurement with a sample free of haemoglobin and five crystalline human haemoglobin (Sigma-Aldrich Cat nr H7379) dilutions: 0.05, 0.1, 0.2, 0.4 and 0.6 mg haemoglobin mL^{-1} . Absorbance of each dilution was measured twice. Reference curves always had a regression coefficient superior to 0.99. Results were then expressed in milligrams of leghaemoglobin per gram of fresh nodules ($\mathrm{mg\,g}^{-1}$).

NIR spectra acquisition and preprocessing

NIR spectra of dried nodules were acquired with an NIR hyperspectral line scan camera combined with a conveyor belt (BurgerMetrics SIA, Riga, Latvia) installed at the Walloon Agricultural Research Centre (CRA-W, Belgium). The device is described in detail in Vermeulen *et al.* ¹⁸ and Eylenbosch *et al.* ¹⁹ Acquisitions were performed with HyperPro VB software (BurgerMetrics SIA, Riga, Latvia). Before image acquisition, the system was calibrated with a white ceramic plate (white reference) and by blocking the entrance of reflected light (dark reference). Bad

pixels were detected and removed at the same time. A dark reference was automatically performed before the acquisition of each image. One hyperspectral image was acquired for each nodule sample (one sample per growing pot) and all nodules used for the study were scanned on the same day.

Nodule samples were put on a conveyor belt progressing at $1200\,\mu\text{m}\,\text{s}^{-1}$ under the camera. For each pixel, one spectrum of 209 wavelengths was acquired in the $1118-2424\,\text{nm}$ range. Pixel resolution was $0.31\,\text{mm}$ wide. One mean spectrum representative of each nodule sample was then computed from each hyperspectral image with the HyperSee software (BurgerMetrics SIA, Riga, Latvia). A threshold was used to segment images and remove background prior to mean spectra computation. The mean spectrum of each individual nodule was also computed.

In order to remove noisy regions at the beginning and the end of the NIR spectra, only the wavelengths between 1432 nm and 2368 nm were kept. ¹⁹ The remaining noise and the increasing trend observed in the mean spectra were removed by use of Savitzky–Golay smoothing (order 0, filter width 7) and detrend transformations, respectively. Spectra were also normalised with standard normal variate (SNV) transformation (Figure 1).

Linking NIR hyperspectral and multispectral imaging to leghaemoglobin content

For each modality (harvest date \times fertilisation), the leghaemoglobin content and the average NIR spectrum

from one nodule sample were kept for validation of the regression model. This sample was randomly selected among the four replicates. The remaining data were used for model calibration (two samples were unusable due to handling errors). In total 22 mean spectra were used for calibration and 8 others for validation.

PLS regression model construction

PLS regression was used to build a calibration model between the leghaemoglobin content and the NIR spectra of pea nodules. The number of latent variables was chosen in order to reduce the cross-validation classification error average. A 10-group venetian blind cross-validation was performed.

This work was performed using the PLS Toolbox 7.8 software (Eigenvector Research, Inc., Wenatchee, WA, USA) working with Matlab R2015a software (The Math Works, Inc., Natick, MA, USA).

Optimal wavelength selection and multiple linear regression fitting

Spectra acquired with hyperspectral imaging contain a large number of wavelengths. Most of the wavelengths are, therefore, likely to provide redundant information and make little or no contribution to the predictions. Identification of wavelengths carrying the maximum spectral information would allow the simplification of the model and further technology transfer to more simple

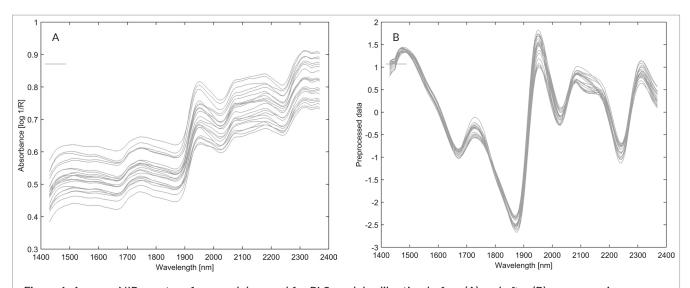


Figure 1. Average NIR spectra of pea nodules used for PLS model calibration before (A) and after (B) preprocessing (Savitzky–Golay smoothing (order 0, filter width 7), detrend and SNV transformations).

acquisition systems, allowing for the implementation of multispectral imaging.

The standard coefficients (β -coefficients) of the PLS regression were used to analyse and identify which wavelengths were the most significant in explaining the variations observed in leghaemoglobin content. ²⁰ As two proximal wavelengths might contain redundant information, rather than identifying the wavelength with the highest score (as a positive or negative value), it was decided to identify the peaks in the graphs of β -coefficients against wavelengths (cf. Results section). The wavelengths for which peaks were identified were then used in a multiple linear regression model to predict leghaemoglobin content.

Algorithms used to select the wavelengths and create the multiple linear regression model were developed using Matlab R2015a software (The Math Works, Inc., Natick, MA, USA).

Statistical analysis

Statistical analyses of nodule numbers, leghaemoglobin content and dry aerial biomass were performed with R software.²¹ Analysis of variance (ANOVA) and pairwise comparisons with Student–Newman–Keuls tests were made using the agricolae package.²²

Three criteria were used to evaluate the quality of the models and their ability to predict leghaemoglobin content: root mean square error (RMSE), determination coefficient (R^2) and the ratio of standard deviation to RMSE called ratio of prediction to deviation (RPD).^{23,24} According to Saeys *et al.*,²⁴ model predictions can be considered as good when the R^2 value is between 0.82 and 0.90. They are classified as good or excellent when the RPD values are between 2.5 and 3.0 or above 3.0, respectively.

Results and discussion

Pea aerial biomass and nodule development

At each harvest date, the number of nodules on each plant root system and the aerial biomass were measured. For the statistical analysis, these measurements were taken per pot (three plants). Both variables were significantly affected by accumulated heat units (*p*-value < 0.01, Table 1). The ANOVA reported no significant impact of nitrogen fertilisation (results not shown). Throughout the study, nodules of different ages and sizes were observed. An increase in nodule number was observed between the first and the second harvest; after which the number of nodules tended to stabilise. Aerial dry biomass increased with an exponential shape with accumulated heat units. At the end of the study, most developed plants had 13 leaves.

The leghaemoglobin content measured in the harvested nodules with the cyanmethaemoglobin method was comprised between $1.4\,\mathrm{mg\,g^{-1}}$ and $4.2\,\mathrm{mg\,g^{-1}}$ (Table 1 and Figure 2). These concentrations were consistent with those reported in previous studies. ^{1,4} A broad range of leghaemoglobin content in nodules was therefore observed, as expected with the protocol established.

The analysis of leghaemoglobin content in pea nodules showed a highly significant effect (*p*-value < 0.001, Table 1) of harvest time. An increase in leghaemoglobin content was observed between the first and third harvest. No statistical differences were reported between the last two harvest dates. Contrary to our expectations, there was no observable effect of nitrogen fertilisation on leghaemoglobin content in nodules (result not shown). This absence of effect could be due to the high amount of nitrogen already present in the growth substrate, which was taken from a field where peas had been grown in

Table 1. Number of nodules, aerial dry biomass production (g) and nodule leghaemoglobin content (mg g⁻¹ fresh nodule) measured per growing pot (three plants) at the four harvest dates. Mean and standard deviation are presented for each harvest date. N-fertilised and non-fertilised plants are pooled. Mean values followed by the same letter are not significantly different (Student-Newman-Keuls, α = 0.05).

Accumulated	Nodules		Aerial dry biomass		Leghaemoglobin content	
heat units (°C-days)	Mean (g pot ⁻¹)	SD (g pot ⁻¹)	Mean (g pot ⁻¹)	SD (gpot ⁻¹)	Mean (mg g ⁻¹)	SD (mg g ⁻¹)
500	95.6 (b)	27.4	0.79 (c)	0.22	1.63 (c)	0.25
630	179.7 (a)	89.9	1.22 (bc)	0.63	2.50 (b)	0.56
760	218.3 (a)	72.5	2.03 (b)	0.75	3.42 (a)	0.48
820	216.0 (a)	58.3	4.69 (a)	1.52	3.52 (a)	0.43

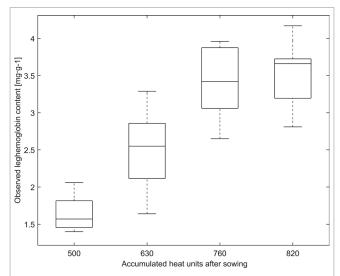


Figure 2. Leghaemoglobin content (mg leghaemoglobin g⁻¹ fresh nodule) measured with the cyanmethaemoglobin method at each harvest time (expressed in heat units (°C-days) accumulated after plant sowing).

the previous crop season. Furthermore, the growing conditions, i.e. higher temperature of the substrate due to the cultivation performed in pots within the greenhouse, have likely increased soil mineralisation which also increased the nutrient content available for the plants. Figure 3 presents the leghaemoglobin content for the different harvest dates, gathering together fertilised and unfertilised datasets.

PLS regression

The main goal of this work was to investigate whether leghaemoglobin content could be predicted from average NIR spectra of pea nodules acquired by NIR hyperspectral imaging. The first attempt was made using a PLS regression to link leghaemoglobin content, measured with the cyanmethaemoglobin method, and NIR spectra of the nodule. In the calibration phase, 22 nodules samples were used. Five latent variables were used to predict the leghaemoglobin content and explained 99.98% of leghaemoglobin variance. The first latent variable explained 98.51% of the variance.

The PLS regression showed a determination coefficient (R^2) of 0.87 and a root mean square error of calibration (RMSEC) of 0.32 (Figure 3). The cross-validation of the PLS model gave an R^2 of 0.74 and a root mean square error (RMSECV) of 0.45. Lower values obtained with cross-validation were probably due to the small number of samples.

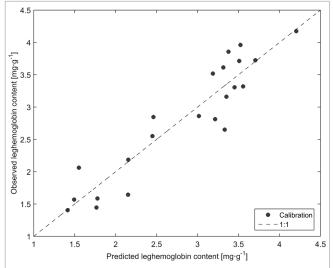


Figure 3. Calibration results of the PLS regression model calibrated on leghaemoglobin content measured with cyanmethaemoglobin method and predicted leghaemoglobin content of nodules on the basis of their NIR spectra. Results are expressed in mg leghaemoglobin g^{-1} fresh nodules. The regression was constructed on five latent variables. Leghaemoglobin was predicted with a RMSEC of 0.32 and a determination coefficient (R^2) of 0.87.

The validation of the PLS model on eight samples gave an R^2 of 0.90 and a root mean square error of prediction (RMSEP) of 0.27 (Figure 4). The RPD was 3.42. Predictions were therefore classified as good.²⁴ The leghaemoglobin contents of the samples used for the validation were well predicted by the regression, and the predictions obtained from these samples were better than those obtained with cross-validation: the R² was higher and the RMSEP was lower than the RMSECV. These samples were not completely independent from those used for the regression model calibration because they were obtained under the same conditions. However, the division of samples, or spectra in a dataset, acquired under the same conditions into calibration and validation sets is common in chemometric studies to estimate the potential of a methodology. 23,25-27

Validation on single nodules

As stated in the Materials and method section, NIR hyperspectral imaging allows one spectrum per pixel to be acquired. The results obtained with the PLS model (PLS regression section) were based on the computation of an average spectrum representative of the whole sampling, i.e. a sample including all

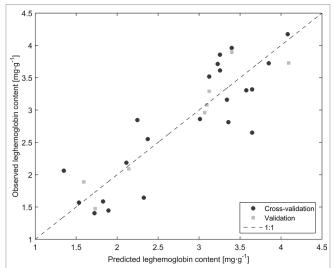


Figure 4. Cross-validation (dark grey circles) and validation (light grey squares) results of the PLS regression model. Leghaemoglobin content was measured with cyanmethaemoglobin method and predicted on the basis of nodule NIR spectra. Results are expressed in mg leghaemoglobin $\rm g^{-1}$ fresh nodules. The regression was constructed on five latent variables. Leghaemoglobin was predicted with a RMSECV of 0.45 and a determination coefficient (R^2) of 0.74. When the regression was applied to the spectra used for the validation, the RMSEP was 0.27 and the R^2 was 0.90.

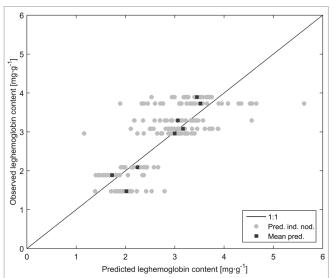


Figure 5. Predicted values of leghaemoglobin on individual nodules (mg leghaemoglobin g⁻¹ fresh nodule). Predictions were made on each mean NIR spectrum computed for each nodule (light grey circles). Nodules were contained in height samples (validation dataset). The mean leghaemoglobin content (dark grey squares) computed on the whole sample was linked with the reference value of leghaemoglobin (observed leghaemoglobin content) measured for the sample with the cyanmethaemoglobin method.

nodules harvested for each growing pot. To evaluate the variability within each sampling, which would make it possible to evaluate the leghaemoglobin content of individual nodules separately, we computed one mean spectrum per nodule within each image (using the information contained in all pixels related to each nodule of the image). The average pixel number for one nodule was 37 (ranging between 10 and 290 pixels per nodule). The PLS model was then applied to each mean spectrum representative of each individual nodule of the validation dataset. The results are presented in Figure 5.

The variability observed in leghaemoglobin content predicted from mean spectra tended to be higher where the measured content was greater, i.e. in nodule samples harvested from the oldest plants. These plants had nodules of different ages and diameters on their roots. Our observations were consistent with results reported for soybean by Sato *et al.*, ¹⁶ who showed that plants have nodules of different diameters and that the leghaemoglobin content increases with the nodules' diameter.

Multiple linear regression model applied to a restricted number of wavelengths

Figure 6 reports the β -coefficients corresponding to each wavelength when the PLS analysis was performed (here, the Matlab function *plsregress* was used to retrieve the coefficients). The graph corresponded to the results obtained from plotting the output of the PLS regression using the first component of the PLS obtained with the *plsregress* function. The percentage of variance explained in this case was about 71%. As stated in the Materials and method section, rather than identifying the highest coefficients, it was decided to identify the peaks within the β -coefficients graph (Figure 6).

The wavelengths identified were: 1461, 1537, 1732, 1934, 2022, 2110 and 2236 nm (Figure 7). A multiple linear regression model was calibrated with the absorbance values measured at these wavelengths to predict leghaemoglobin content. The calibration displayed similar performances to those observed in the complete PLS approach, with an R^2 of 0.85 and an RMSEC of 0.35.

During the validation phase, the results again revealed similar performances, with an R^2 of 0.86 and an RMSEP

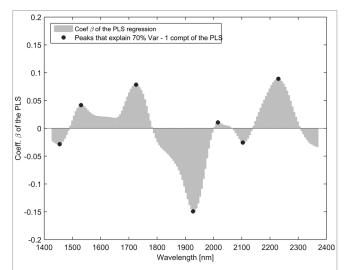


Figure 6. Most relevant wavelengths selected using the graphical representation of coefficients against the corresponding wavelength.

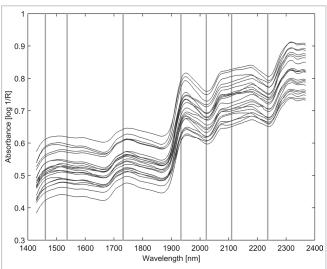


Figure 7. Identification (vertical grey lines) of the seven wavelengths (1461, 1537, 1732, 1934, 2022, 2110 and 2236 nm) that were used to perform multiple linear regression analysis. The wavelengths of interest were reported on the graphical representation of the pea nodule spectra used to calibrate the multiple linear regression model.

of 0.33. Graphical representations were fairly similar to those reported when applying the complete PLS approach.

Not surprisingly, most of the identified wavelengths were linked to chemical structure containing carbon and nitrogen. These wavelengths are situated in the first overtone, the second overtone and the combination band regions of the NIR spectrum.²⁸ Key wave-

lengths identified at 1461, 1934 and 2110 nm are located in the regions of N-H bonds. The 1934 nm wavelength is also linked to moisture. Wavelengths identified at 1732 nm and 2236 nm are located in the C-H bond regions.

Conclusion

This paper has proposed a new method for leghaemoglobin quantification in dried pea nodules using NIR hyperspectral imaging combined with chemometrics as an alternative to the classical reference cyanmethaemoglobin method. The PLS regression gave good results when the whole spectrum of pea nodules was used.

This PLS regression applied to the mean spectrum of individual nodules showed consistent results: nodules collected from the same plant had different leghaemoglobin content, due to the size and age of nodules, and their mean leghaemoglobin content as well as the variability of their leghaemoglobin content tended to increase with plant age.

Using the peaks identified in the β -coefficients graph resulting from the PLS analysis applied to the calibration dataset, seven wavelengths of interest were chosen. A multiple linear regression model was calibrated on these wavelengths to predict leghaemoglobin content. This simpler model gave similar results on the validation dataset to those of the PLS regression applied to the whole spectrum.

Both proposed approaches showed good correlation with the reference method. However, they show some advantages over the reference approach: (i) they are quicker; (ii) they are simpler (once the model is calibrated, the quantification does not depend on operator ability); (iii) they do not involve the use of chemical and potentially dangerous reagents; (iv) the leghaemoglobin content of a single nodule can be predicted. Considering all these advantages and the first results acquired in this study, NIR hyperspectral imaging and multispectral analysis have demonstrated strong potential for evaluating the activity of pea nodules through the quantification of leghaemoglobin content. They appear suited to be used within large-scale field studies in order to improve knowledge of nitrogen fixation and protein accumulation in legume crops.

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