

Quantification of protein in wheat using near infrared hyperspectral imaging: Performance comparison with conventional near infrared spectroscopy

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Abstract

Hyperspectral imaging is a powerful technique that combines the advantages of near infrared spectroscopy and imaging technologies. Most hyperspectral imaging studies focus on qualitative analysis, but there is growing interest in using such technique for the quantitative analysis of agro-food products in order to use them as universal tools. The overall objective of this study was to compare the performance of a hyperspectral imaging instrument with a classical near infrared instrument for predicting chemical composition. The determination of the protein content of wheat flour was selected as example. Spectra acquisition was made in individual sealed cells using two classical near infrared instruments (NIR-DS and NIR-Perstop) and a near infrared hyperspectral line-scan camera (NIR-HSI). In the latter, they were also acquired in open cells in order to study the possibility of accelerating the measurement process. Calibration models were developed using partial least squares for the full wavelength range of each individual instrument and for the common range between instruments (1120–2424 nm). The partial least squares models were validated using the “leave-one-out” cross-validation procedure and an independent validation set. The results showed that the NIR-HSI system worked as well as the classical near infrared spectrometers when a common wavelength range was used, with an r^2 of 0.99 for all instruments and Root Mean Square Error in Prediction (RMSEP) values of 0.15% for NIR-HSI and NIR-DS and 0.16% for NIR-Perstop. The high residual predictive deviation values obtained (8.08 for NIR-DS, 7.92 for NIR-HSI, and 7.56 for NIR-Perstop) demonstrate the precision of the models built. In addition, the prediction performance with open cells was almost identical to that obtained with sealed cells.

Keywords

Hyperspectral imaging, near infrared spectroscopy, wheat, protein content

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Introduction

Since near infrared (NIR) spectroscopy was first applied in agriculture by K.H. Norris in 1964, to measure moisture in grains,¹ this technique has been widely used by the agro-food industry for determining the chemical composition and other quality properties in biological samples. NIR spectroscopy is based on sample absorption at specific wavelengths of incident radiation. This absorption depends on the chemical composition and physical state of the sample. Like other vibrational spectroscopic techniques, NIR is classified as an indirect analytical technique because a calibration step is required in order to subsequently predict a particular property from the spectra of unknown samples. Robust and flexible NIR

instruments, as well as easy-to-use software for building calibration models, are now available for routine control. Chemical determination based on NIR spectra is fairly simple, rapid, nondestructive, and cost effective. It requires a low input of reagents and can be implemented at-line, on-line, or in-line at the site of food

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or feed production. At the laboratory level, under ISO17025 conditions and taking account of all the sample preparation and analysis steps, up to 100 samples of agro-food products can be analyzed in duplicate in one day and by one analyst.²⁻⁴

More recently, NIR hyperspectral imaging (HSI) has been developed as a powerful technique that combines the advantages of NIR spectroscopy and imaging technologies. The great advantage of HSI is its ability to acquire simultaneously spectral and spatial information from a sample. Whereas classical NIR instruments collect spectral data from an area of a fraction of the sample, HSI collects spatially distributed NIR spectra responses at many subsample areas or pixels (usually hundreds or thousands) of the sample.⁵⁻⁷ Each hyperspectral image is represented in a 3D spectral cube, usually called hypercube, with 2D spatial information and 1D spectral information (i.e. absorbance at specific wavelengths). The most important HSI applications in the agro-food industry are in detecting defects,^{8,9} discriminating botanical species, cultivars and quality classes,¹⁰⁻¹² determining fruit ripeness⁷⁻¹³ and chemical composition,¹⁴⁻¹⁶ and detecting and quantifying contaminants.^{6,17-24}

Traditionally, hyperspectral instruments are classified into three groups, depending on the way the hypercube is generated: point-scan, line-scan, and plane-scan instruments. Point-scan instruments successively acquire a spectrum at a simple spatial location of the scene of interest. For this, a spectrometer (equipped with a single detector), coupled with a microscope equipped with an automatic sample stage, is used to perform the mapping in order to construct the hypercube. Line-scan instruments, through the use of a 2D focal plane array (FPA) detector, allow the spectra of several pixels from a line to be collected simultaneously. The hypercube is generated by moving the camera or samples horizontally in order to scan the whole scene. Plane-scan instruments allow the absorbance at continuous wavelengths of the scene of interest to be gathered successively. With this equipment, an FPA detector is also used, but a device is needed to select the absorbance at specific frequency, the hypercube being built through a step-by-step acquisition of the absorbance images at the successive wavelengths. In agro-food product analysis, line-scan instruments seem to be the best compromise for speed of analysis, flexibility, and possible use for on-line control.^{10,25}

As noted earlier, NIR HSI is used mainly in qualitative studies and its use in quantification is still limited. However, there is an interest for the development of global tools where bulk calibration of classical quantitative constituents for quality control could be determined with the same apparatus used for the determination of properties that are easy or could only be detected with an imaging system mainly related to qualitative aspects as contaminant or fraud detection. In this direction, only in recent years, several studies have demonstrated the feasibility of using an HSI system for the

quantitative prediction of the internal composition of agro-food products. Several studies have reported on the use of HSI to predict, inter alia, the content of water, fat, protein, and total saturated and total unsaturated fatty acids in red meat²⁶⁻²⁹; fat and protein content in cheese³⁰; moisture and fat content in various species of fish³¹; soluble solids, moisture content, and acidity in fruits and vegetables³²⁻³⁵; and moisture, fat, starch, and oleic acid content in cereals.^{14,36,37} Studies have also been conducted to predict the content of minor compounds such as anthocyanins and flavonols in grape skin³⁸ and grape seed,³⁹ respectively; total pigment in red meat⁴⁰; synthetic astaxanthin coating and total volatile basic nitrogen in fish^{41,42}; and total glucosinolate in freeze-dried broccoli.⁴³ Macronutrients in oilseed rape leaves⁴⁴ and wheat leaves⁴⁵ have been also determined using HSI. In addition to predicting internal composition, NIR HSI has been used for predicting water holding capacity, color, pH, tenderness, and microbiological attributes, such as total viable counts of bacteria, in red meat^{46,47} and in fish,⁴⁸⁻⁵² as well as firmness in fruits and vegetables.^{7,34}

Based on these references, HSI appears to be a powerful technique for the quantitative prediction of the composition of food products. However, it is not known how similar its performance is to any other classical NIR instrument. Knowing more about its comparative performance would enhance the potential of HSI for quantitative aspects until now mainly conducted with classical NIR spectrometers. New HSI instruments are now cheaper and faster, and the development of adequate chemometric tools and computer software will undoubtedly help to optimize the exploration of both spatial and spectral information for qualitative and/or quantitative determination. With the decreasing cost of HSI instruments, there is growing interest in finding out if they can be used in the same way as classical NIR spectrometers for determining quantitative parameters. So far as we know, there have been only a few studies comparing HSI with conventional NIR spectroscopy for quantitative prediction. They have focused on agro-food products usually considered a challenge for classical NIR techniques because of the heterogeneous physical nature of the sample (raw instead of ground material) and the heterogeneous spatial distribution of the parameters of interest. Burger and Geladi³⁰ compared an HSI instrument based on plane-scan technology with two NIR spectrophotometers (i.e. a scanning grating one with rotating sample holders and an FT-NIR one equipped with a fiber-optic sampling probe) for predicting the protein and fat content of cheese. By selecting a subset region of interest in the hyperspectral images and reducing it to a single average spectrum, the authors obtained almost identical results to those obtained with classical NIR instruments. The prediction error in terms of the average bias of the HSI instrument was 0.6% for protein and 0.2% for fat content compared with those obtained with the classical NIR

instruments. The scanning grating spectrometer showed the best results (0.2% for protein content and 0.5% for fat content) and the FT-NIR the worst results (3.0% for protein content and 0.7% for fat content). Xing et al.⁵³ compared a line-scan HSI system with an FT-NIR spectrophotometer for predicting alpha-amylase activity in individual wheat kernels. The performance of the HSI system when using the average spectra from the region of interest (i.e. the germ region, where the alpha-amylase is mainly located) was almost identical to the performance achieved with classical FT-NIR instruments. For the HSI system, the best model produced a coefficient of determination r^2 for the validation set of 0.82 and a root mean square error (RMSE) of 0.54. The predictive ability of the FT-NIR was lower; the best model was obtained with the original data (r^2 and RMSE were 0.71 and 11.76, respectively). Mendoza et al.,⁵⁴ however, compared the ability of an on-line hyperspectral scattering system and a short NIR spectrometer (USB400, with a range of 460–1100 nm) for predicting the firmness and soluble solids content (SSC) in three apple cultivars over two seasons. Based on SEP and residual predictive deviation (RPD) values, the NIR instrument was always better at predicting SSC, although the RPD values were < 2.5. In contrast, the hyperspectral scattering system was better at predicting firmness. Ignat et al.⁵⁵ showed that an HSI system based on electronically tuned band-pass filter (AOTF) performed poorly compared with a short NIR spectrometer (USB2000, with 350–1000 nm of spectral range) and a Liga SWIR spectrophotometer in predicting several quality parameters in intact bell peppers. The best prediction models were found for the SSC and the dry matter content. For SSC, the RPD values were 3.9, 3.3, and 2.6 for the short NIR, Liga SWIR and HSI systems, respectively; for the dry matter, the RPD values were 3.8, 3.0, and 2.4, respectively. Rady et al.⁵⁶ compared the potential of an NIR hyperspectral line-scan system with two classical NIR spectrometers (i.e. an NIR transmittance system and a visible/NIR interactance system using a fiber optic), inter alia, for predicting the glucose and sucrose content and the SSC of two potato cultivars. All the instruments performed well for glucose, particularly sliced samples. The visible/NIR interactance system showed the best prediction models in terms of the correlation coefficient r^2 and RPD (≥ 0.90 and 2.14, respectively) for glucose. For the NIR hyperspectral system, the best model for glucose was an r^2 and RPD of 0.74 (r^2 of 0.55) and 1.49, respectively.

This study sought to assess the potential of an NIR hyperspectral line-scan system for predicting some important quality parameters of agro-food matrices and comparing its performance with that of two classical NIR instruments widely used in the agro-food industry and research laboratories. The overall objective was to determine if an HSI instrument is as efficient for predicting chemical composition as a classical NIR instrument. This is performed because there is an

increasing need of developing global instruments, i.e. instruments for the simultaneous determination of different properties. On the basis of this consideration, it could be more worthy to calibrate protein on wheat kernels, for instance, where simultaneously the same analysis could reveal the presence of pests, contaminants, etc. The collection of spectra on a very fine (pixel-based) spatial level, the heterogeneity of the imaged substance as well as size and shape characteristics for intact objects can be obtained for defects/contaminants detection; while with the average of pixels of the scene or fraction of it to obtain a representative spectrum can be of utility to perform quantitative analysis.

In this study, the determination of the protein content of wheat flour, historically one of the most common uses of NIR spectroscopy in the agricultural and food industry,⁵⁷ was selected as an example. Wheat flour samples are easy to handle using classical NIR because the grinding process homogenizes the product and protein content is a parameter that is stable over time. Moreover, cereals products are usually studied with a HSI in order to detect their purity, homogeneity, and contamination among others.^{58–60}

Materials and methods

Samples

A set of 79 wheat flour samples was used for this study. The samples were collected in 2013, with 57 coming from Belgian mills and 22 from the Bureau Interprofessionnel des Etudes Analytiques (Bipea) in France, an independent association that provides testing programs to laboratories.

Reference analysis

The Dumas combustion method^{61,62} was used to determine the total nitrogen, and therefore the crude protein, content. The method involves the total combustion of samples under oxygen. Via subsequent oxidation and reduction tubes, nitrogen is quantitatively converted to N₂. Results are given as % or mg nitrogen, which may be converted into protein by using conversion factors.⁶³ The reference value (% of protein content) for the Bipea samples was 11.06 ± 0.90 , with a minimum of 9.50 and a maximum of 12.67. For the rest of the samples, the mean value was 11.10 ± 1.28 and varied between 9.00 and 14.40. The estimated uncertainty of the method for determination of total protein content is of 0.27.

Instrumentation and spectra acquisition

Two classical NIR instruments working in reflection mode, as well as a NIR hyperspectral line-scan camera were used in this study. Table 1 gives the main characteristics of the instruments used.

The first NIR instruments is a NIR System DS2500 (Foss—hereafter referred to as NIR-DS) that operates

Table 1. Characteristics of NIR instruments and NIR-HSI: total wavelength range, common wavelength range, spot size, and analysis time by sample.

Instrument name	Wavelength range (nm)	Common range (nm)	Spot size ^a (cm ²)	Analysis time ^b (s/sample)
NIR-DS	400–2498	1120–2424	1.8	66.0
NIR-Perstop	1100–2498	1120–2424	1.8	57.6
NIR-HSI	1118–2425	1120–2424	3.0	28.8

NIR: near infrared.

^aFor NIR-DS and NIR-Perstop, spot size is fixed at 15 cm in diameter; for NIR HSI, a total of 2601 spectra (51 pixels × 51 pixels) were selected from the center of each image.

^bDetermined from the total time required by instrument for the measurements of all samples (n = 79).

in reflection mode in the 400–2498 nm range, with a spectral resolution of 0.5 nm. It averages a certain number of spectra collected at different locations in a sample cup (rotating sampling device) during analysis. Spectra were collected using the ISIScan Nova software across the original wavelength range. Each spectrum was the average of 32 scans performed on the sample and it was acquired in 66 s. The sample area measured was about 2 cm².

The second NIR instrument is a 5000 Autocup DVP6BX instrument (Foss—hereafter referred to as NIR-Perstop) that operates in reflection mode in the 1100–2498 nm range, with a spectral resolution of 0.5 nm. It also has a rotating sampling device.

Spectra were collected using ISIScan software across the original wavelength range and each spectrum was the average of 32 scans performed on the sample. The sample area measured was about 2 cm², and 57 s was needed to analyze a cell. The difference of time of analysis (i.e. 9 s) between both NIR instruments (i.e. NIR-DS and NIR-Perstop) was due mainly to the auto-sampler device of the NIR-Perstop instrument.

The third instrument is a NIR hyperspectral line-scan camera (Burgermetrics—hereafter referred to as NIR-HSI) or push-broom imaging system was used to collect hyperspectral images. The NIR-HSI instrument was a SWIR XEVA CL 2.5 320 TE4 camera using a spectrograph. It has a cooled, temperature-stabilized Mercury–Cadmium–Telluride detector combined with a conveyor belt that presents the sample to the camera. There is more information on this instrument in Vermeulen et al.²³ and Dale et al.¹⁰

Spectra acquisition was made using HyperPro software (Burger-Metrics SIA, Riga, Latvia). All the images consisted of 420 lines of 320 pixels acquired at 209 wavelength channels (1128–2425 nm) and 32 scans per image. A 15 mm lens was used to analyze the total width of the plate. The lens was set up to cover 10 cm of the width of the conveyor belt. The conveyor belt speed was fixed at 3 mm/s (i.e. 20 lines/s) so as to produce clear images. In order to compensate for offset due to the dark current, the light source temperature drift, and

the lack of spatial lighting uniformity, the spectral imaging system was calibrated with a dark image (by blocking the lens entrance) and a white image (background) collected from a standard white reference board (empty Teflon plate). The spectra were then automatically corrected.

After spectral acquisition, a total of 2601 spectra (51 × 51 pixels) corresponding to a sample area of ±3 cm² were selected from the center of each image and averaged in order to obtain the mean spectra of each sample measured. In the configuration selected, the time required to analyze one sample was 28.8 s.

For each instrument, the 79 samples of wheat flour were measured in individual sealed cells with a glass cover, first with the NIR-DS instrument, then with the NIR-HSI instrument, and finally with the NIR-Perstop instrument. Sealed cells were used to prevent small moisture content changes in the samples and to ensure that the same portion of each sample was presented in the same condition to the different instruments. In order to study whether or not the measuring process in the line-scan system could be accelerated, after the NIR-Perstop measurements with sealed cells, spectra were also acquired in open cells and compared with spectra of the sealed cells. In all cases, the samples were measured randomly, although always in the same order for all the instruments. The NIR measurements were done in duplicate (i.e. the batch of samples was measured consecutively two times according to the protocol described before). The analysis of all the samples with the three instruments was simultaneously performed over three consecutive days.

Figure 1 shows the mean spectra of the 79 wheat flour samples obtained with each instrument (i.e. NIR-DS, NIR-Perstop, and NIR-HSI). For all instruments, typical reflectance spectra are obtained with the traditional bands and shape observed for ground wheat.⁶⁴ The different bands are related mainly to the different stretching vibrational bands assigned to the O–H, N–H, and C–H groups.⁶⁵

Spectral data analysis

The spectral data were processed using Matlab software, v. 7.0 (The Mathworks, Inc, Natick, MA, USA). Chemometric tools were applied using the PLS Toolbox (v. 4.11, Eigenvector, Inc., Manson, WA, USA).

Principal component analysis (PCA) was used for exploratory analysis in order to extract the maximum amount of information from each set of data. Calibration models were developed using partial least squares (PLS) for predicting the protein content for the three instruments. These models were built using the full wavelength range, which varied from instrument to instrument (see Table 1) and also with the common range between them (reduced wavelength range), which corresponded to 1120–2424 nm. The number of PLS latent variables has been determined through the ‘leave-one-out’ cross-validation (LOOCV) procedure,

and then an external/independent validation has been applied. For LOOCV, one sample was left out and the multivariate models were constructed with the rest of the samples. The process was repeated until all the samples had been used once in the validation set. For independent validation, the samples were split into two groups using two strategies: a calibration model with 75% of samples randomly selected and validation with the remaining 25% (**Strategy 1**); and a calibration model with the first 60 samples in order of measurement, and validation with the remaining 19 samples

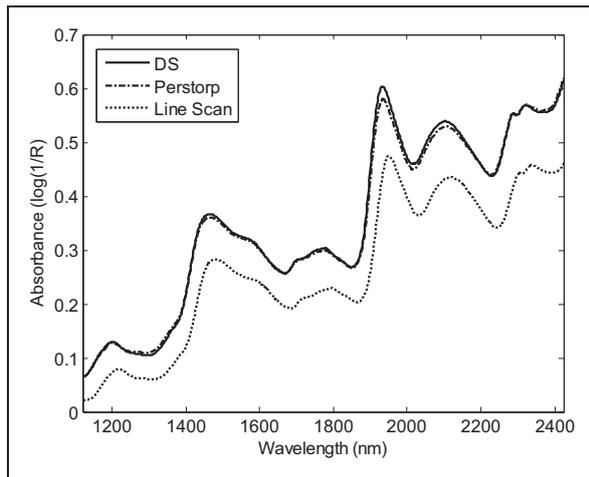


Figure 1. Mean spectrum of wheat flour obtained with the NIRS-DS, NIRS-Perstop and NIR-HSI instruments.

(**Strategy 2**). The aim of Strategy 2 was to check stability over time, as mentioned earlier. In both strategies, the samples of the validation set were independent (i.e. they came from different sources, wheat cultivars, growing conditions, harvesting regimes, etc.).

The quality of the constructed models was evaluated using the RMSE for calibration (RMSEC), cross-validation (RMSECV), and external validation (RMSEP). The coefficients for calibration and prediction (r^2) were determined, as well as the RPD for calibration and validation ($RDP_{cal}=SD/RMSEC$ and $RDP_{val}=SD/RMSEP$, respectively; with SD =standard deviation). The best prediction models were those with higher values of r^2 and RPD and lower values of RMSEC and RMSECV. Values of RPD greater than 3.0 indicate excellent prediction accuracy.⁶⁶

Results and discussion

The PCA score and loading plots of the sealed samples are shown in Figure 2. The first two PCs explained 99.73% of the total variation in the raw NIR spectral matrix. PC1 showed a clear separation between the spectra from all the instruments clearly visible in the loading plot and PC2 allowed the NIR spectra from the NIR-HSI instrument to be discriminated from those coming from the NIR-DS and NIR-Perstop instruments.

The Hotelling's T^2 , a measure of the distance from the multivariate mean to the projection of each sample

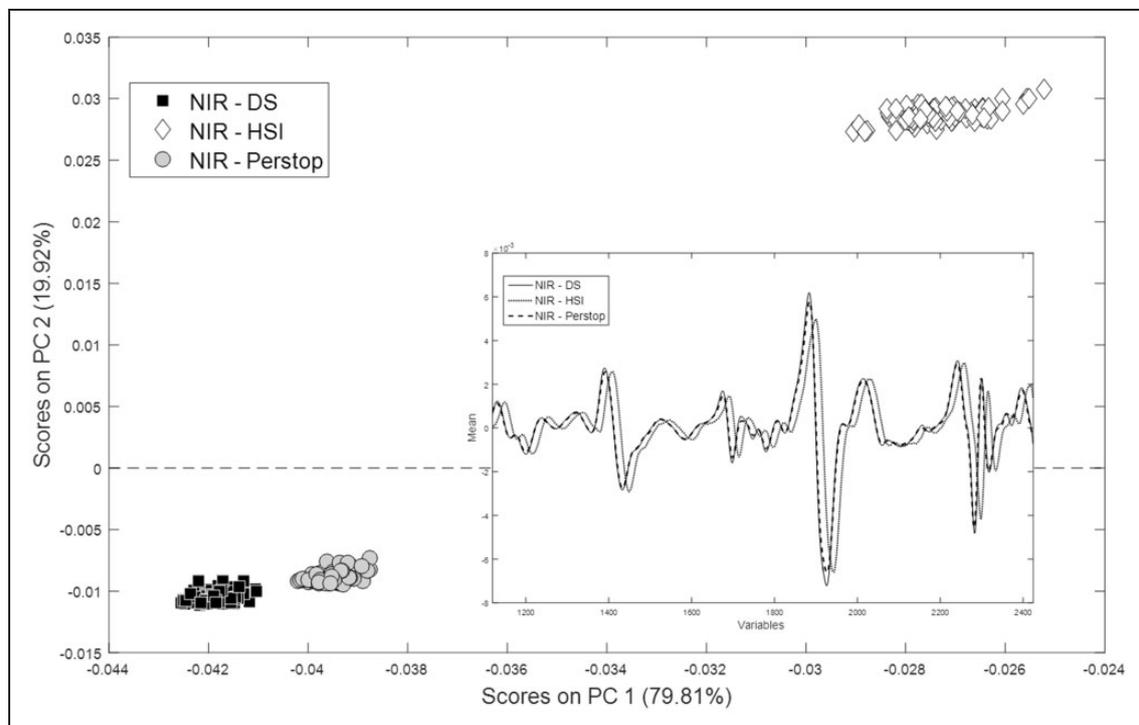


Figure 2. Principal Component Analysis (PC1 and PC2).

onto these two PCs, gave average values of 7.62, 6.87, and 15.38 for NIR-DS, NIR-Perstop, and NIR-HSI, respectively, indicating that the distance between the conventional NIR instruments was smaller, and so they are almost equivalent. These results were confirmed when a three-nearest-neighbors distance calculation was performed, with average values of 0.08, 0.05, and 0.179 for NIR-DS, NIR-Perstop, and NIR-HSI, respectively. This suggests that there was more variability in the raw data coming probably from HSI due to a large measurement surface and non-homogeneous light distribution.

Table 2 shows the results for the PLS models for the determination of protein content for all the instruments tested and using both the whole wavelength ranges and the common range. The most efficient preprocessing tool involved the use of standard normal variate followed by a second derivative Savitzky–Golay (window: 11, polynomial: 2). As shown in the table, good predictions with sealed cells were obtained when considering the total wavelength range with an r^2 of 0.99 for all the instruments. NIR-DS gave the best results in terms of RMSEP (0.14%) compared to NIR-Perstop (0.16%) and NIR-HSI (0.17%). The generally good performance of the NIR spectroscopy instruments (NIR-DS and NIR-Perstop) was not really a surprise, as NIR spectroscopy is used routinely in the cereal industry worldwide for quantifying protein content and other chemical components of wheat flour.⁶⁷ However, the similar performance of NIR-HSI demonstrates the potential of this technique for similar quantitative applications. Looking at the RPD values, which were higher than 3.0, the independent validations produced excellent predictions for protein, regardless of instrument.

When the common wavelength range (1120–2424 nm) was considered, Strategy 1 gave the best prediction model, with a similar r^2 (0.99) for all instruments. NIR-HSI showed the same RMSEP values (0.15%) as NIR-DS. NIR-Perstop showed the highest value (0.16%) and an almost identical performance for the total wavelength. The RPD values (8.08, 7.92, and 7.56 for NIR-DS, NIR-HSI, and NIR-Perstop, respectively) showed excellent predictions of protein content. These results demonstrate the potential of NIR-HSI compared with conventional NIR instruments for quantification, agreeing with earlier studies that showed even better prediction of protein and fat content in cheese by hyperspectral plane-scan imaging than FT-NIR,³⁰ better prediction of alpha-amylase activity in individual wheat kernels by hyperspectral line-scan imaging than FT,⁵³ and better prediction of firmness in apple fruits using an on-line hyperspectral scattering system than with using a short NIR spectrometer.⁵⁴

The results differed, however, with regard to the outputs of Strategy 2 (i.e. the first measured 60 samples of the wheat flour set were used for calibration and the remaining 19 samples for validation). Here, the performances of the NIR-DS and NIR-Perstop

Table 2. PLS models for the total and common wavelength range.

	Total wavelength range						Common wavelength range Strategy 1 ^a						Common wavelength range Strategy 2 ^b											
	LV	R^2	RMSEC (%)	RMSECV (%)	RPDcal	r^2	RMSEP (%)	RPDval	LV	R^2	RMSEC (%)	RMSECV (%)	RPDcal	r^2	RMSEP (%)	RPDval	LV	R^2	RMSEC (%)	RMSECV (%)	RPDcal	r^2	RMSEP (%)	RPDval
NIR-DS	8	0.96	0.15	0.22	5.27	0.99	0.14	8.55	6	0.97	0.16	0.20	5.84	0.99	0.15	8.08	6	0.97	0.15	0.20	5.87	0.98	0.16	7.56
NIR-Perstop	6	0.95	0.20	0.25	4.72	0.99	0.16	7.56	6	0.95	0.20	0.24	4.86	0.99	0.16	7.56	5	0.96	0.21	0.25	4.65	0.98	0.14	8.68
NIR-HSI ^b	7	0.97	0.15	0.19	6.18	0.99	0.17	7.07	7	0.96	0.14	0.23	5.09	0.99	0.15	7.92	7	0.98	0.11	0.20	5.96	0.94	0.23	5.15

LV: Latent Variables; PLS: partial least squares; RMSEC: RMSE for cross-validation; RMSECV: RMSE for cross-validation; RMSEP: Root Mean Square Error in Prediction for an external set; RPD: residual predictive deviation.

^aStrategy 1: Calibration model with 75% of samples randomly selected and validation with the remaining 25%; Strategy 2: Calibration model with the first 60 samples according to the order of measurement and validation with the remaining 19 samples.

^bSamples were measured in sealed cells on the NIR-DS instrument, followed by the NIR-HSI instrument, and finally by the NIR-Perstop instrument.

instruments were similar, but worse than in Strategy 1. For both instruments, the r^2 was 0.98, although the RMSEP was higher with NIR-DS (0.16%) than with NIR-Perstop (0.14%). For NIR-HSI, the r^2 was 0.94, the RMSEP 0.23, and the RPD fell by 35% compared with Strategy 1. Although the mean RPD value was higher than 3.0, denoting excellent prediction accuracy, there was an important loss in prediction ability. The results of Strategy 2 therefore indicated a decline in the stability of the NIR-HSI instrument over time, which is undesirable for routine analysis at industrial or laboratory level. This could be easily solved by calibrating the instrument several times a day and before each analysis of a set of samples.⁶⁸

Figure 3 shows the prediction of the calibration and validation sets for the total wavelength range using Strategy 1 and the prediction of the validation set for the reduced/common wavelength range using Strategies 1 and 2.

In addition to the study on sealed cells, a tentative HSI study was done on open cells (not shown in table). The prediction performance for the total wavelength range in open cells (r^2 of 0.98 and RMSEP of 0.20%) was almost identical to that obtained with the HSI

instrument in sealed cells (r^2 of 0.99 and RMSEP of 0.17%). The RPD value fell by 17% for the sealed cells, indicating excellent accuracy for protein content prediction. Given that the time needed for analyzing open cells fell by about 50% with the NIR-HSI instrument compared to the other instruments (Table 1), this is an interesting factor to take into account. The use of open cells can therefore increase the speed of measurement with NIR-HSI and could also be accelerated by simultaneous measurement of two cells; hence the time required would fall by up to about 14 s/measurement.

Conclusion

To date, the use of HSI has been based mainly on its capacity for discrimination in qualitative studies, such as those focusing on fraud or contaminant detection. In our study, the comparison with classical NIR instrumentation demonstrated the potential of NIR-HSI for quantifying the chemical composition of the samples.

In particular, the study showed that when a common wavelength range is used for all the instruments and the same sample sets, a hyperspectral line-scan system worked as well as a classical NIR spectrometer and

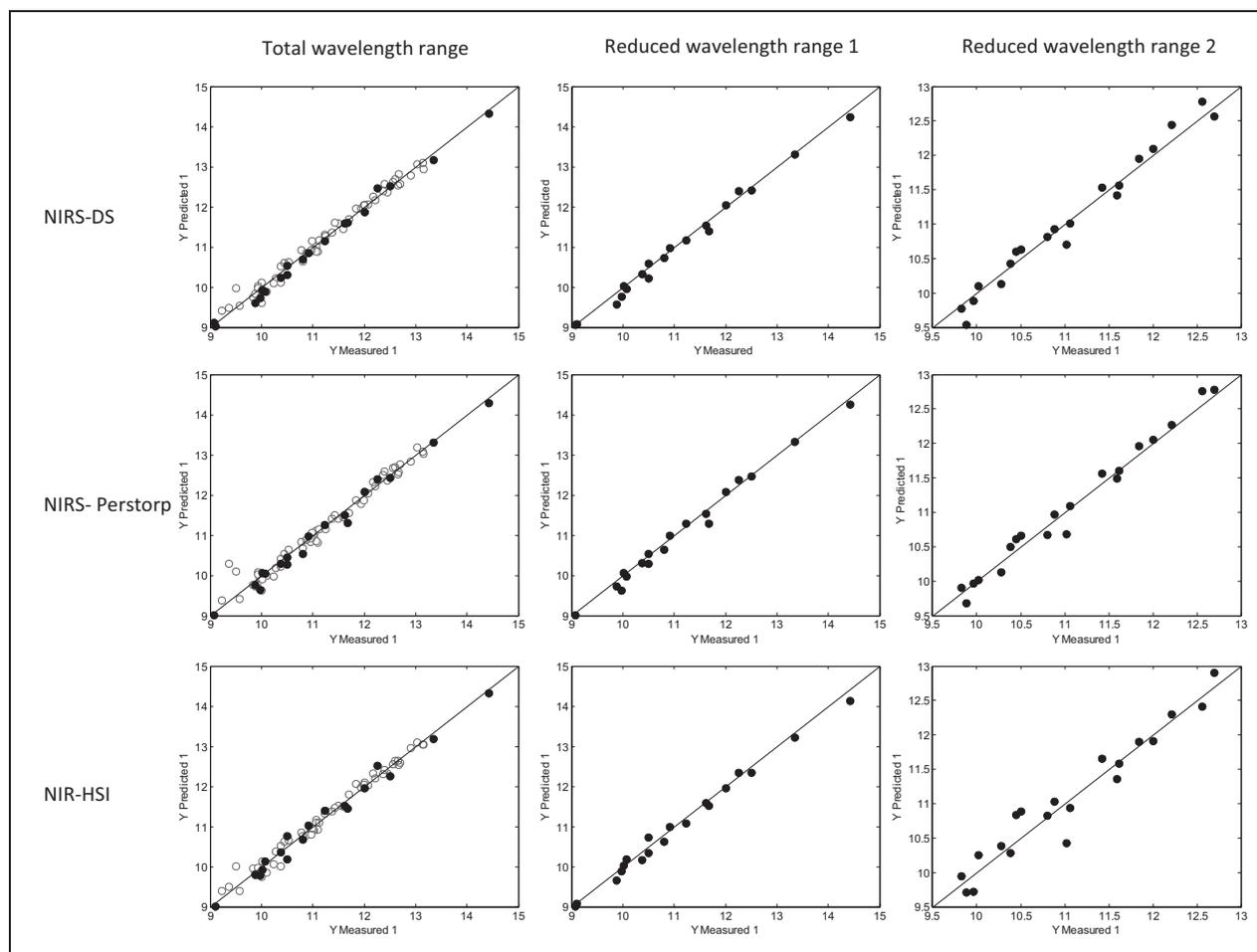


Figure 3. NIR predicted data versus reference data for protein content, for total wavelength range (strategy 1) and common wavelength range (strategy 1 and 2).

the time required for analyzing a sample decrease by at least half. In addition, it showed that using open cells increased the number of samples that could be analyzed over time, paving the way to the development of a complete methodology for on-line analysis.

The study demonstrated, then, the feasibility of using NIR-HSI for predicting the protein content of wheat flour and highlighted the potential of this technology for the cereal industry, given its already demonstrated ability for sorting wheat into different classes according to growing conditions, visible or internal defects, and contaminants.

The main drawback of this type of system is (i) the loss of stability over time, which affects the performance of hyperspectral systems in quantitative studies, but this can be solved by carrying out the appropriate calibration strategy; and (ii) its price, which is probably the main limitation of using this technique for quantification. However, this could be solved as looking at this technique as a multitool instrument, i.e. a system that can be used as a classical NIR spectrometer for quantitative quality estimations, as proved in this study, and at the same time an instrument that gives a fast and reliable solution for the qualitative detection of abnormal ingredients or products, often demonstrated in the scientific literature.

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