



SLOVENSKI STANDARD

SIST EN 15948:2015

01-junij-2015

Nadomešča:
SIST EN 15948:2012

Žito - Določevanje vlage in beljakovin - Metoda z uporabo bližnje infrardeče spektroskopije v celih zrnih

Cereals - Determination of moisture and protein - Method using Near-Infrared-Spectroscopy in whole kernels

Getreide - Bestimmung der Feuchte und des Proteins - Verfahren der Nahinfrarot-Spektroskopie bei ganzen Körnern

Céréales - Détermination de la teneur en eau et en protéines - Méthode utilisant la spectroscopie dans le proche infrarouge sur des grains entiers

Ta slovenski standard je istoveten z: EN 15948:2015

ICS:

67.060	Žita, stročnice in proizvodi iz njih	Cereals, pulses and derived products
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EUROPEAN STANDARD
NORME EUROPÉENNE
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ICS 67.060

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English Version

**Cereals - Determination of moisture and protein - Method using
Near-Infrared-Spectroscopy in whole kernels**

Céréales - Détermination de la teneur en eau et en
protéines - Méthode utilisant la spectroscopie dans le
proche infrarouge sur des grains entiers

Getreide - Bestimmung der Feuchte und des Proteins -
Verfahren der Nahinfrarot-Spektroskopie bei ganzen
Körnern

This European Standard was approved by CEN on 5 January 2015.

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EUROPEAN COMMITTEE FOR STANDARDIZATION
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EN 15948:2015 (E)**Foreword**

This document (EN 15948:2015) has been prepared by Technical Committee CEN/TC 338 "Cereal and cereal products", the secretariat of which is held by AFNOR.

This European Standard shall be given the status of a national standard, either by publication of an identical text or by endorsement, at the latest by October 2015, and conflicting national standards shall be withdrawn at the latest by October 2015.

Attention is drawn to the possibility that some of the elements of this document may be the subject of patent rights. CEN [and/or CENELEC] shall not be held responsible for identifying any or all such patent rights.

This document supersedes EN 15948:2012.

The following modifications were made in this new edition:

- Annexes have been enhanced with information on 2 other NIR instruments based on the results of the interlaboratory tests and the models of prediction.

According to the CEN-CENELEC Internal Regulations, the national standards organizations of the following countries are bound to implement this European Standard: Austria, Belgium, Bulgaria, Croatia, Cyprus, Czech Republic, Denmark, Estonia, Finland, Former Yugoslav Republic of Macedonia, France, Germany, Greece, Hungary, Iceland, Ireland, Italy, Latvia, Lithuania, Luxembourg, Malta, Netherlands, Norway, Poland, Portugal, Romania, Slovakia, Slovenia, Spain, Sweden, Switzerland, Turkey and the United Kingdom.

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1 Scope

This European Standard defines a routine method for the determination of moisture and protein in whole kernels of barley and wheat using a near-infrared spectrophotometer in the constituent ranges:

a) for wheat:

- 1) moisture content minimum range from 8 % to 22 %;
- 2) protein content minimum range from 7 % to 20 %.

b) for barley:

- 1) moisture content minimum range from 8 % to 22 %;
- 2) protein content minimum range from 7 % to 16 %.

This European Standard describes the modalities to be implemented by the supplier (5.3 and 5.4) and the user of the method.

2 Normative references

The following documents, in whole or in part, are normatively referenced in this document and are indispensable for its application. For dated references, only the edition cited applies. For undated references, the latest edition of the referenced document (including any amendments) applies.

EN ISO 12099:2010, *Animal feeding stuffs, cereals and milled cereal products — Guidelines for the application of near infrared spectrometry (ISO 12099:2010)*

ISO 5725-2, *Accuracy (trueness and precision) of measurement methods and results — Part 2: Basic method for the determination of repeatability and reproducibility of a standard measurement method*

3 Terms and definitions

For the purposes of this document, the terms and definitions given in EN ISO 12099:2010 apply.

4 Principle

The method is based on Near-Infrared (NIR) spectroscopy, an indirect, correlative technique to predict the concentration of various constituents in organic samples. Linear or non-linear regression modelling is used to relate NIR spectra to moisture or protein concentrations determined by officially approved standard methods (e.g. artificial neural network - ANN, Partial Least Square Regression - PLS).

5 Method of analysis

5.1 General

According to this document, the method of analysis is defined as the association between a NIR instrument and a model of prediction.

EN 15948:2015 (E)**5.2 Near Infrared Instrument**

Based on diffuse reflectance or transmittance measurement covering the wavelength region of 700 nm to 2 500 nm or segments of this or at selected wavelengths.

5.3 Prediction models

Each model for the prediction of protein and moisture contents in whole grain of wheat and barley is amongst others defined by:

- the number of samples used for the calibration development;
- the constituent ranges covered in the model for moisture and protein;
- the temperature range of the samples;
- the number and performance of involved reference labs;
- the stability of the model i.e. by number of harvests covered;
- the calibration file defined by its name and its IT name (for example CHECKSUM) insuring its integrity;
- the seasonal, geographic and genetic variations covered.

5.4 Initial validation of the model

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5.4.1 General

Since NIR analysis is an indirect, correlative technique, the results shall be validated against chemical analysis reference methods. It is important that the reference methods used are officially approved such as the methods described in the standards previously cited (Clause 2). The purpose of validation is to determine the root mean square error of prediction which depends at the same time on the correlation, the bias and the slope.

The root mean square error between chemical analysis methods and predictions shall be compared to calibration performance specifications and/or historical performance.

5.4.2 Initial validation sample set

The initial validation of a calibration model shall be done in accordance with EN ISO 12099 using independent test sets of wheat and barley samples, originating from different countries and analyzed by the reference methods given in Clause 2.

Requirements for the validation sample set are:

- at least 200 samples coming from 10 countries (20 representative samples min/country) distributed homogeneously over the entire constituent range;
- the part of the range without any reference sample shall not exceed 0,3 %;
- different scans from one sample shall not be considered as different samples;
- seasonal effects over at least a three year period, temperature effects, instrument variation and the variability of reference data shall be included in the set.

5.4.3 Initial validation performances

The results of the initial validation shall at least fulfil the specifications given in Table 1.

Table 1 — NIR performances for the determination of moisture and protein (see also Annex B)

		Moisture Wheat and barley	Protein Wheat	Protein Barley
Overall accuracy expressed as SEP as constituent % w/w		0,24 %	0,27 %	0,27 %
Constituent concentration in the independent validation data set	Min	8 %	7 % d.m.	7 % d.m.
	Max	22 %	20 % d.m.	16 % d.m.

NOTE The minimum performance given in Table 1 includes the variation of reference data as documented by the number of reference labs involved, regional and genetic variations, the number of countries and crop species involved and the robustness over the last five years (see also Annex B).

5.5 Update of calibration model and validation of new model

The prediction model in accordance with this standard shall be updated by the one issuing the calibration model to ensure inclusion of new climatic crop conditions and new varieties introduced on the market. These updates shall be made by keeping the original database with addition of the new samples as needed.

The new prediction model shall be updated according to EN ISO 12099.

Validation shall be made according to the initial validation (5.4) and include at least 20 new samples.

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6 Sampling

Sampling is not part of the method specified in this European Standard. A recommended sampling procedure is given in EN ISO 24333 [5].

It is important that the sample analyzed in routine is truly representative for the batch and has not been damaged or modified.

7 Procedure

7.1 Preparation of the test sample

No specific sample preparation is required.

7.2 Measurement

Follow the instructions of the instrument manufacturer.

7.3 Local validation of the method

Before use, the method shall be validated on an independent test set that is representative of the sample population to be analyzed. For the determination of bias, at least 10 samples are needed; for the determination of Standard Error of Prediction (SEP, see EN ISO 12099:2010, 6.5) at least 20 samples are

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needed. Validation shall be carried out for each sample type, constituent/ parameter and temperature (see EN ISO 12099:2010, 5.4).

Bias or inherent systematic error, as described in EN ISO 12099:2010, Clause 6, is exhibited when the predicted results of a specific sample group or product show a mean offset value when compared to their reference values. This may occur with unique sample types.

The bias (i.e. mean difference between the chemical analysis results and the predicted results) may or may not be statistically significant. Based on the procedure described in EN ISO 12099, a bias confidence limit may be calculated.

When this limit is exceeded, a bias is implemented in the instrument software and the validation process repeated. Refer to the manufacturer instructions and to EN ISO 12099 for procedure.

7.4 Periodical adjustment of the instrument

To ensure its accuracy, each instrument shall be checked at least annually, against the reference method, either directly or through a master instrument.

The execution of this check shall be performed on samples covering a range as wide as possible, taking into account seasonal, geographical and genetic variations.

The number of samples for the adjustment should be sufficient for the statistics used to check the performance. For the determination of the bias, at least 10 samples are needed, for the determination of standard error of prediction (SEP) and for the slope adjustment, at least 20 samples are needed.

7.5 Checking instrument stability

See EN ISO 12099:2010, Clause 9.

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7.6 Follow up of method performance

Performance of the method shall be checked at least annually, against reference methods to secure the constant adequacy of the model with the requirements of this standard (see 5.4.2).

This performance test shall be made on samples selected from the pool of analyzed samples. It may be necessary to resort to some sampling strategy to ensure a balanced sample distribution over the entire calibration range and to ensure that samples with a commercially important range are covered. At least 20 samples are needed (to expect a normal distribution of variance).

For instruments operated in a network and adjusted against a master instrument, it is sufficient to run the performance check of the method of this last one.

The adjustment (7.4) respecting the requirements of this clause may be used for the follow-up of the method performance.

It is recommended to participate in an internationally accepted proficiency testing scheme (PTS) that includes NIRS predicted results and results generated by following the standards specified in Clause 2.

8 Calculation and expression of results

The software of the instrument calculates the results for moisture and protein and displays them in % w/w (g/100 g) to two decimal places.

If multiple measurements are made on the same sample, calculate the arithmetic mean.

Express final results to two decimal places.

9 Accuracy and precision of the method

9.1 Accuracy

The accuracy of the prediction model is determined by validation in accordance with EN ISO 12099 and expressed by the Standard Error of Prediction (see EN ISO 12099:2010, Table 1). The Standard Error of Prediction (SEP) is an expression of the bias corrected average difference between predicted and reference values predicted by the model when applied to a set of samples not included in the derivation of the model. The values also include the uncertainty of reference results.

The predicted results will not in more than 5 % of cases deviate more than $1,96 \times \text{SEP}$ (as determined in the paragraph above) from the best estimate of the true value.

NOTE As NIR is an indirect method, the typical standard deviation of reproducibility for the used reference methods are given here for comparison:

- moisture (EN ISO 712) = 0,16 %;
- protein (EN ISO 20483) = 0,20 %;
- protein (EN ISO 5983-2) = 0,20 %;
- protein (CEN ISO/TS 16634-2) = 0,21-0,26 %.

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9.2 Precision

9.2.1 General

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The precision of the prediction model shall be determined from an interlaboratory test organized according to ISO 5725-2 and at least fulfil the performance criteria of repeatability and reproducibility given below.

Details of an example of an interlaboratory test are summarized in Annex A. The precision data given below are derived from this example.

Figure A.1 and Figure A.2 show that the repeatability and the reproducibility are independent of the concentration. The figures in Annex B show that the dispersion is identical over the validated range (Figure B.1 and Figure B.2). Therefore the model can be used in the whole validated range, even though the interlaboratory trial covered a smaller range.

9.2.2 Repeatability

The absolute difference between two independent single test results, obtained using the same method on identical test material in the same laboratory by the same operator using the same equipment within a short interval of time will not in more than 5 % of cases be greater than the repeatability limit r ($r = s_r \times 2,8$) with:

$$r_{\text{protein}} = 0,42 \%$$

$$r_{\text{moisture}} = 0,15 \%$$

9.2.3 Reproducibility

The absolute difference between two single test results, obtained using the same method on identical test material in different laboratories with different operators using different equipment, will not in more than 5 % of cases be greater than the reproducibility limit R ($R = s_R \times 2,8$) with:

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$$R_{\text{protein}} = 0,45 \%$$

$$R_{\text{moisture}} = 0,25 \%$$

9.2.4 Critical difference

9.2.4.1 General

When the difference between two averaged values obtained from two test results under repeatability or reproducibility conditions shall be assessed, the repeatability or reproducibility limit cannot be used, one shall use the Critical Difference (CD).

9.2.4.2 Comparison of two groups of measurements in one laboratory

The critical difference (CD) between two averaged values obtained from two test results under repeatability conditions is equal to:

$$CD = 2,8 s_r \sqrt{\frac{1}{2n_1} + \frac{1}{2n_2}} = 2,77 s_r \sqrt{\frac{1}{2}} = 1,98 S_r$$

where

s_r is the standard deviation of repeatability;

n_1 and n_2 are the numbers of test results corresponding to each of the averaged values;

CD_r (protein) = 0,30;

CD_r (moisture) = 0,11.

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9.2.4.3 Comparison of two groups of measurements in two laboratories

The critical difference (CD) between two averaged values obtained in two different laboratories from two test results under repeatability conditions is equal to:

$$CD = 2,8 \sqrt{s_R^2 - s_r^2 \left(1 - \frac{1}{2n_1} - \frac{1}{2n_2} \right)} = 2,8 \sqrt{s_R^2 - 0,5 s_r^2}$$

where

s_r is the standard deviation of repeatability;

s_R is the standard deviation of reproducibility;

n_1 and n_2 are the number of test results corresponding to each of the averaged values;

CD_R (protein) = 0,32;

CD_R (moisture) = 0,23.

10 Test Report

The test report shall specify:

- a) all information necessary for the complete identification of the sample;
- b) the sampling method used (if known);
- c) the application model and instrument used with reference to this European Standard;
- d) all operating details not specified in this European Standard, or regarded as optional, together with details of any incidents which may have influenced the test result(s);
- e) the test result(s) obtained.

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