

# Prediction of the Heating Value of Biomass Fuel from IR and NIR Data



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## 1. Motivation

**Biomass** is becoming increasingly important as a **renewable source of energy**. Apart from traditional incineration of wood, recent developments also consider the incineration of cereals.

**Spectroscopy** is a promising alternative to time-consuming calorimetric experiments for determining an important property of a fuel: its **heating value**.

This study aims at developing **PLS calibration models** for the prediction of higher heating values of **wood and cereal** samples, based on infrared and near-infrared data.

## 2. Experimental

### 2.1 Samples

20 wood samples: sawdust of spruce, pine and larch with varying amounts of additives (bark, rye/maize flour, starch)  
15 cereal samples: wheat, rye, barley, maize and triticale flours

### 2.2 Instrumentation

Four different set-ups for spectroscopic measurements:

**A.** NIR-spectrometer with sapphire ball probe  
(Brimrose Luminar 5030)

**B.** NIR-spectrometer with rotating sample cup  
(Brimrose Luminar 5030)

**C.** VIS-NIR-spectrometer with Direct Contact Analyser  
(Foss NIRSystems 6500)

**D.** IR-spectrometer with ATR-Durascope sampler  
(Bruker Equinox 55)

### 2.3 Data

**NIR absorbance data (A. - C.):** 1100-2300 nm,  $\Delta\lambda = 8$  nm, 1<sup>st</sup> derivative Savitzky-Golay, resulting in  $p = 145$  features

**IR absorbance data (D.):** 4000-600  $\text{cm}^{-1}$ ,  $\Delta\nu = 8$   $\text{cm}^{-1}$ , 1<sup>st</sup> derivative Savitzky-Golay, resulting in  $p = 435$  features

**Heating values:** The higher heating value in kJ/kg dry biomass is determined by the bomb calorimetric method [1], range: 18,143-19,125 kJ/kg

## 3. Chemometrics

### 3.1 Variable Selection

Variable selection by Genetic Algorithm (GA) using software MobyDigs [2]

- regression method: OLS
- performance criterion (fitness):  $R_{adj}^2$ , adjusted squared correlation coefficient between  $y$  and  $\hat{y}$  for full cross validation [3]
- max. number of features selected in a model: 15
- number of iterations per trial: 1-1.4 million
- typical computation time: 60-120 minutes
- use variables present in the 10 best models

### 3.2 Repeated Double Cross Validation for prediction performance of PLS Models

Application of new software developed in R [4,5].

- The data set is randomly partitioned into  $s = 4$  segments: 3 segments for calibration, 1 segment as test set.
- A PLS model is derived from the calibration set with optimum number of PLS components estimated by cross-validation.
- Application of PLS model to test set results in  $n/s$  predicted values  $\hat{y}_i$ .
- Systematic variation gives a  $\hat{y}$  for each object.
- The whole process is repeated  $k = 10$  times.
- Finally,  $k \cdot n = 10 \cdot 35 = 350$  values  $\hat{y}$  are available.

Performance criteria derived from prediction errors:

$SEP_{test}$  standard deviation of prediction errors

$TI_{90}$  tolerance interval of prediction errors (difference of 95 % and 5 % percentile)

$PC_{med}$  median of  $s-k$  calculated optimum numbers of PLS components

### 3.3 Full Cross Validation for prediction performance of PLS Models

For reference, PLS regression as implemented in software Unscrambler [6] has been performed.

$SEP_{CV}$  standard deviation of prediction errors by full cross validation (leave-one-out)

$PC_{opt}$  optimum number of PLS components given by software

## 4. Results

**Table 1:** Prediction performance of PLS models using all features of NIR (A.-C.) and IR data (D.)

data set	$p$	$SEP_{test}$	$PC_{med}$	$SEP_{CV}$	$PC_{opt}$
A.	145	141	1	124	4
B.	145	115	6	102	5
C.	145	89	5	87	8
D.	435	141	3	137	2

**Table 2:** Prediction performance of PLS models using GA selected features of NIR (A.-C.) and IR data (D.)

data set	$p$	$SEP_{test}$	$PC_{med}$	$SEP_{CV}$	$PC_{opt}$
A.	19	96	14	62	14
B.	21	94	10	56	17
C.	20	96	11	70	10
D.	19	48	13	62	7

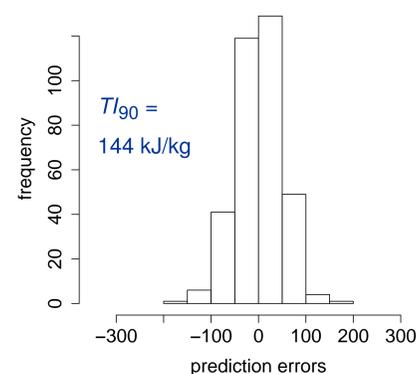
$p$  number of features

$PC$  number of PLS components (*med.* median, *opt.* optimum)

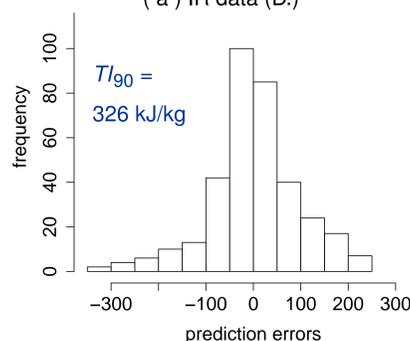
$SEP$  standard deviation of prediction errors (in kJ/kg)

$SEP_{test}$  test sets in repeated double cross validation (10 repetitions)

$SEP_{CV}$  leave-one-out cross validation

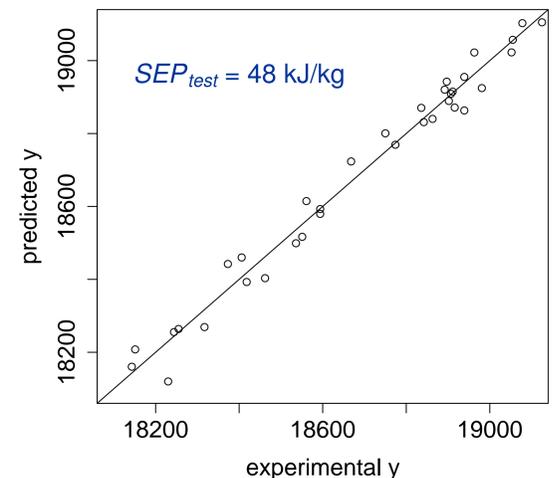


(a) IR data (D.)



(b) NIR data (A.)

**Figure 1:** Frequency distribution and tolerance interval of prediction errors of PLS models using 19 features of (a) IR data (best model) and (b) NIR data



**Figure 2:** Prediction of higher heating values (kJ/kg) by a PLS model using 19 IR absorbances selected by a genetic algorithm. The predicted values are means from 10 replicates of double cross validation.

## 5. Conclusions

- Easily available (Near-)Infrared spectroscopy data are very promising for the determination of heating values of biomass.
- Variable selection by Genetic Algorithm improves prediction performance in most cases, especially for IR data.
- Differences in prediction quality due to instrumental set-up (NIR data) decrease after variable selection.
- A single PLS-model can be used for the heating value of wood as well as cereal samples.
- Simple leave-one-out cross validation tends to give too optimistic results, which can be misleading for variable selection.
- Repeated double cross validation allows realistic estimation of prediction performance.
- Implementation of repeated double cross validation in software R is fast with typical computation times of 10 seconds per job.

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## 7. References

1. Friedl, A., Padouvas, E., Rotter, H., Varmuza, K., *Anal. Chim. Acta* 544 (2005) 191-198.
2. Software MobyDigs, v 1.0. Talete srl, www.talete.mi.it, Milan, Italy, 2004.
3. Frank, I. E., Todeschini, R., *The data analysis handbook*. Elsevier, Amsterdam, 1994.
4. Software R, v 2.5.1. R Development Core Team, www.r-project.org, 2007.
5. Mevik, B. H., Wehrens, R.: *J. Statistical Software* 18 (2007), issue 2.
6. Software The Unscrambler v 9.0. Camo Process AS, www.camo.no, Oslo, Norway, 2004.