## Introduction

# A New Variable Selection Method Based on All Subsets Regression

Andreas Liebminger<sup>1</sup>, Leonhard Seyfang<sup>2</sup> Peter Filzmoser<sup>2</sup>, Kurt Varmuza<sup>1</sup>\*

<sup>1</sup> Institute of Chemical Engineering
<sup>2</sup> Institute of Statistics and Probability Theory
Vienna University of Technology, Austria

\* Presenting author Laboratory for Chemometrics, Institute of Chemical Engineering, Vienna University of Technology Getreidemarkt 9/166, A-1060 Vienna, Austria kvarmuza@email.tuwien.ac.at, www.lcm.tuwien.ac.at

Poster Presentation: 10th SSC 2007, Scandinavian Symposium on Chemometrics 11 - 15 June 2007, Lappeenranta, Finland An ideal variable selection method for regression models would find one or more subsets of variables which have optimum prediction performance.

Usually,

- not prediction performance is optimized during variable selection;
- no exhaustive test of all possible variable subsets is possible;
- empirical variable selection methods have to be applied that are not optimal.

Consequently, the prediction performance of regression models - obtained from different variable subsets - has to be estimated separately.

#### This study

- presents the new variable selection method FASS, by combining forward selection with fast all subsets regression;
- compares FASS with other variable selection methods (for instance a genetic algorithm);
- applies a "repeated double cross validation" for estimating the prediction performance of PLS regression models.

#### Forward selection combined with All SubSets Regression Typical parameters (software in R) [1,2]



# Strategy and Data

# All subsets regression (FASS)

Exhaustive treatment of all variable subsets up to 31 variables. Function "regsubsets" in package "leaps" in R [2]; typ. computation time 2 s per run, 1 - 60 minutes in a FASS application. Regression method OLS; performance criterion *BIC* (Bayesian information criterion, Schwarz criterion, *SIC*), similar to Akaike criterion [3, 11],

 $BIC = \ln(RSS/n) + k \ln(n) / n$  for normally distributed residuals

*n*, no. of objects; *p*, no. of variables + 1 (for intercept); *RSS*, sum of squared prediction errors. *BIC* penalizes a large number of variables.

#### Genetic algorithm (GA)

Software MobyDigs [4]. Regression method: OLS; performance criterion (fitness): adjusted squared correlation coefficient,  $_{ADJ}R^2$ , between  $\gamma$  and  $\hat{\gamma}$  for full cross validation [3]. Maximum number of selected variables is 15, typical computation time 30 - 120 minutes.

#### Prediction performance of PLS models

**Repeated double cross validation.** The data set is randomly partitioned into *s* (typ. 4) segments. A calibration set consists of *s* -1 segments, the remaining segment is a test set. A PLS model is derived from the calibration set (cross validation is used to estimate the optimum number of PLS components), and is applied to the test set, resulting in n/s predicted values  $\hat{y_i}$ . Systematic variation gives a  $\hat{y}$  for each object. The whole process is repeated *k* times (typ. 10-100). Finally, *k.n* predicted values are available. From the prediction errors several performance criteria are derived, e.g.: *SEP*<sub>TEST</sub> (standard deviation), difference of 95% and 5% percentile (confidence interval), density distribution (for visual inspection). Typically, 10 variable subsets (from different selection methods) have been tested by this repeated double cross validation. New software in R [2, 5]; typ. comp. time 2 minutes. Leave-one-out cross validation using all data (Unscrambler [6]).

### Results

#### Data sets

- **OXY:** n = 180, p = 57. Concentration change of isotope <sup>18</sup>O in precipitation ( $\gamma$ ) modeled by meteorological and geographical variables [7].
- **PAC:** n = 209, p = 467. GC-retention indices ( $\gamma$ ) of polycyclic aromatic compounds [8], modeled by molecular descriptors (Dragon [9]).
- **TOX:** n = 846, p = 681. GC-retention indices ( $\gamma$ ) of compounds relevant in forensics [10], modeled by molecular descriptors (Dragon [9]).

Dataset	p	Variable selection	<b>SEP</b> TEST	SEP CV
<b>OXY</b> <i>y</i> = (-16.5)-(-5.5)	57 11 15 13	no FASS GA stepwise	1.01 0.83 0.84 1.09	0.90 0.74 0.79 0.77
<b>PAC</b> <i>y</i> = 197-504	467 27 15 22	no FASS GA stepwise	11.0 5.2 7.2 18.7	7.3 5.0 6.7 5.3
<b>TOX</b> <i>y</i> = 1110-3870	681 31 15 33	no FASS GA stepwise	108 82 100 205	80 80 97 74

*n*, number of objects; *p*, number of variables;

SEP, standard deviation of prediction errors (standard error of prediction);

TEST, test sets in repeated double cross validation (10 repetitions);

CV, leave-one-out cross validation

#### Results

Dataset	п	p	Computing time per job [minutes] <sup>1</sup>		
			FASS <sup>2</sup>	GA <sup>3</sup>	
OXY	180	57	0.2	26	
PAC	209	467	7	40	
ТОХ	846	681	180	120	

<sup>1</sup> PC processor AMD Athlon 2.2 GHz; <sup>2</sup> Until no further improvement obtained; <sup>3</sup> Termination after 200000 iterations

 Variable selection by FASS or GA improved prediction performance; stepwise selection was not successful.

- FASS results are similar to GA results or better.
- Advantages of FASS are: less computation time, selection of up to 31 variables (GA in used software allows only 15), more strictly defined algorithm.
- Simple leave-one-out cross validation can be very misleading. A careful estimation of prediction performance is necessary for evaluation of variable selection methods.

#### References

- [1] Liebminger, A.: PhD Dissertation. Vienna University of Technology, Austria, 2006.
- [2] Software R, 2.2.0. R Development Core Team, www.r-project.org, 2005.
- [3] Frank, I. E., Todeschini, R.: The data analysis handbook. Elsevier, Amsterdam, 1994.
- [4] Software MobyDigs, 1.0. Talete srl, www.talete.mi.it, Milan, Italy, 2004.
- [5] Mevik, B.H., Wehrens, R.: J. Statistical Software 18 (2007) issue 2.
- [6] Software The Unscrambler, 9.0. Camo Process AS, www.camo.no, Oslo, Norway, 2004.
- [7] Liebminger, A., Papesch, W., Haberhauer, G., Varmuza, K.: Chemom. Intell. Lab. Syst. (2007), doi: 10.1016/j.chemolab.2007.04.005.
- [8] Lee, M.L., Vassilaros, D.L., White, C.M., Novotny M.: Anal. Chem. 51 (1979) 768.
- [9] Software Dragon, 5.0, Talete srl, www.talete.mi.it, Milan, Italy (2004).
- [10] Garkani-Nejad, Z., Karlovits, M., Demuth, W., Stimpfl, T., Vycudilik, W., Jalali-Heravi, M., Varmuza, K.: J. Chromatogr. A 1028 (2004) 287.
- [11] Box, G.E.P., Jenkins, G.M., Reinsel, G.C.: Time series analysis: Forecasting and control, 3rd ed., Prentice Hall, Upper Saddle River, NJ, 1994.