

# Evaluation of variable selection approaches for pixel-based analysis of GC×GC data

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## Aim

- Study the effect of gas condensate chemical composition on **reactor coke formation during steam cracking**.
- Evaluate **variable (feature) selection** methodologies prior to partial least squares regression (PLSR).

## Introduction

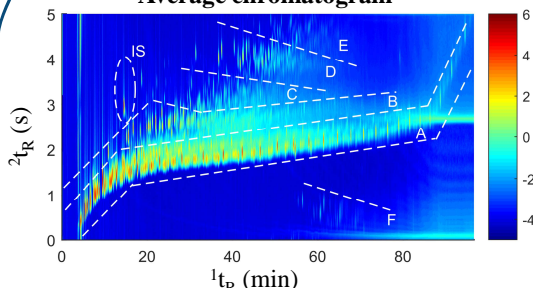
- Pixel-based analysis** enables integration free interpretation of GC×GC data. Therefore all information is retained and analysis is swift.<sup>1</sup>
- PLSR** can be used to correlate information in chromatograms with *e.g.* petroleum properties.<sup>2</sup>
- Variable selection** is needed to remove redundant information and to avoid focusing on large peaks in chromatograms which might not be important.<sup>3</sup>

## GC×GC

- 50 m dimethyl polysiloxane column (RTX-1 PONA, 0.25 mm I.D., 0.5 µm film thickness)
- 2 m phenyl polysilphenylene-siloxane (BPX50, 0.15 mm I.D., 0.15 µm film thickness)
- Dual-stage cryogenic (liquid CO<sub>2</sub>) modulator
- Flame ionization detection
- 8 gas condensates with duplicate analysis and QC samples

## Results and discussion

### Average chromatogram



Representation of the mean of all chromatograms with a logarithmic transformation to visualize the large span between intensities of various analyte groups.

The group-type separation is categorized by: (A) paraffins and naphthenes, (B) monoaromatics, (C) naphthoaromatics, (D) diaromatics, (E) naphthenodiaromatics, (F) triaromatics. Added 3-Chlorothiophene was used as internal standard (IS).

Several feature selection methodologies were evaluated. The RReliefF algorithm resulted in the lowest root mean square error of the cross-validation (RMSECV).

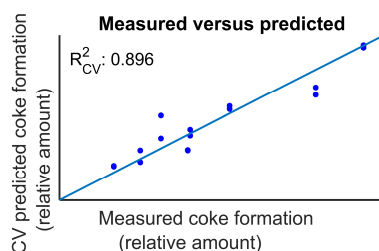
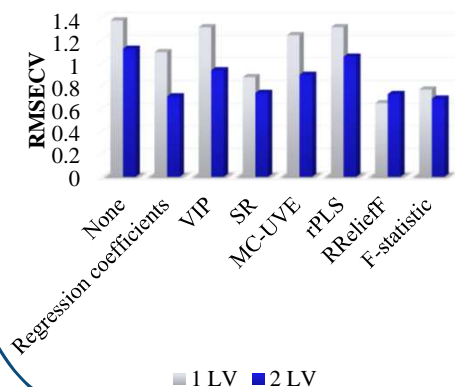
## Signal processing

Alignment	<ul style="list-style-type: none"> <li>Interval correlation optimized warping on unfolded data (1D chromatograms)</li> <li>2D correlation optimized warping</li> </ul>
Baseline correction	<ul style="list-style-type: none"> <li>Minimum value of each modulation</li> </ul>
Normalization	<ul style="list-style-type: none"> <li>Internal standard (3-Chlorothiophene)</li> </ul>
Scaling & transformation (evaluated)	<ul style="list-style-type: none"> <li>None</li> <li>Inverse within-sample standard deviation</li> <li>Base 10 logarithm</li> </ul>

## Conclusions

- Feature selection is a crucial part of the development of PLSR models
- Feature selection indicates important regions in chromatograms
- RReliefF was the most efficient method
- RReliefF offers independent evaluation and does not require an initial PLSR model.
- Reactor coke formation was associated with heavy aromatic compounds and could be used as a predictor

### Comparison of feature selection methods



\*Cross-validation (CV) was performed using leave-one-sample-out, *i.e.* both replicates.

## References

- (1) Furbo, S., et al. (2014). *Anal. Chem.* **86**(15): 7160-7170.
- (2) Pierce, K. M., et al. (2015). *Data Handl. Sci. Techn.* **29**: 427-463.
- (3) Andersen, C. M. and R. Bro (2010). *J. Chemometr.* **24**(11-12): 728-737.

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