

Comparative study of different third-order data generation approaches / HPLC-FD data analysis

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INTRODUCTION

Over the last years, it has been demonstrated that the increase of multi-way data dimensions has a positive impact on analytical figures of merit, e.g. higher sensitivity, lower limits of detection and quantitation, better selectivity, among others. First- and second-order data analyses have become excellent tools for the resolution of complex samples which would result experimentally challenging from the univariate calibration standpoint. On the other hand, even though no additional analytical advantages have been yet proved, third-order data analysis for analytical applications constitutes a field worth to be explored. Although multidimensional instrumental signals are easy to be obtained with the available modern instrumentation, and several chemometric algorithms have been successfully developed to solve multi-way data problems, the way in which the multi-way data are generated may have a significant effect on the final results. In this work, a comparative study of different third-order data generation approaches was carried out. Three methods based on identical liquid chromatographic conditions but coupled to different emission and excitation fluorescence detection system were developed for the analysis of antibiotics in aqueous matrices.

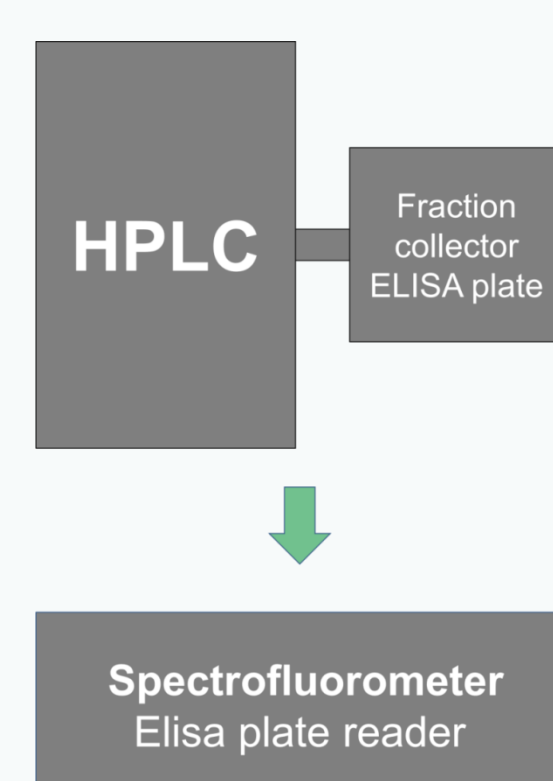
EXPERIMENTAL PROCEDURES

All experiments were performed under same chromatographic conditions, by using an Agilent 1100 LC instrument in isocratic mode.

- Analytical column Zorbax XDB-C18, 75×4.6mm, 3.5µm.

- Mobile phase: HAc buffer pH4.0:ACN:MeOH mixture (71:9:20).
- Column temperature at 35 °C.

1) Collection of fractions



Flow rate: 2 mL min⁻¹

- For each sample:
 - 1 Chromatographic run collecting 25 fractions, every 2 s = 2 min
 - 25 EEMs = 40 min.

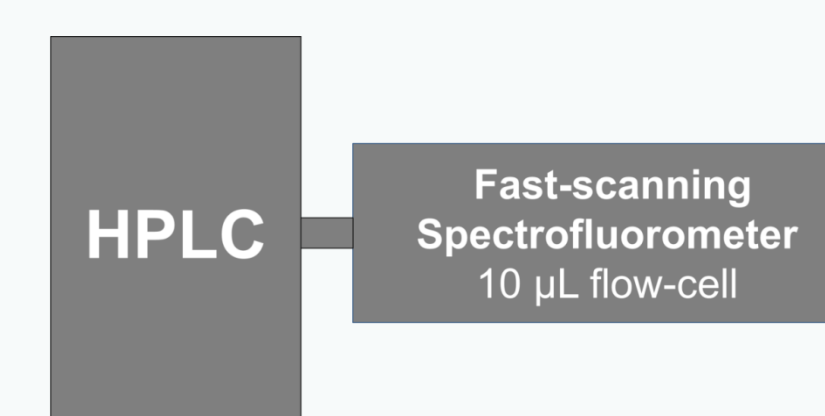
2) Multi-chromatographic run / Multi-excitation



Flow rate: 2 mL min⁻¹

- For each sample:
 - 10 Chromatographic runs recording time-emission matrix at fixed excitation wavelength (different excitation wavelength for each run) = 40 min

3) On-line EEM detection



Flow rate: 0.5 mL min⁻¹

- For each sample:
 - 1 Chromatographic run recording 25 sequential EEMs = 7 min

DATA ANALYSIS

Analysis time per sample

(HPLC + FD)= 45 min

Matrix size

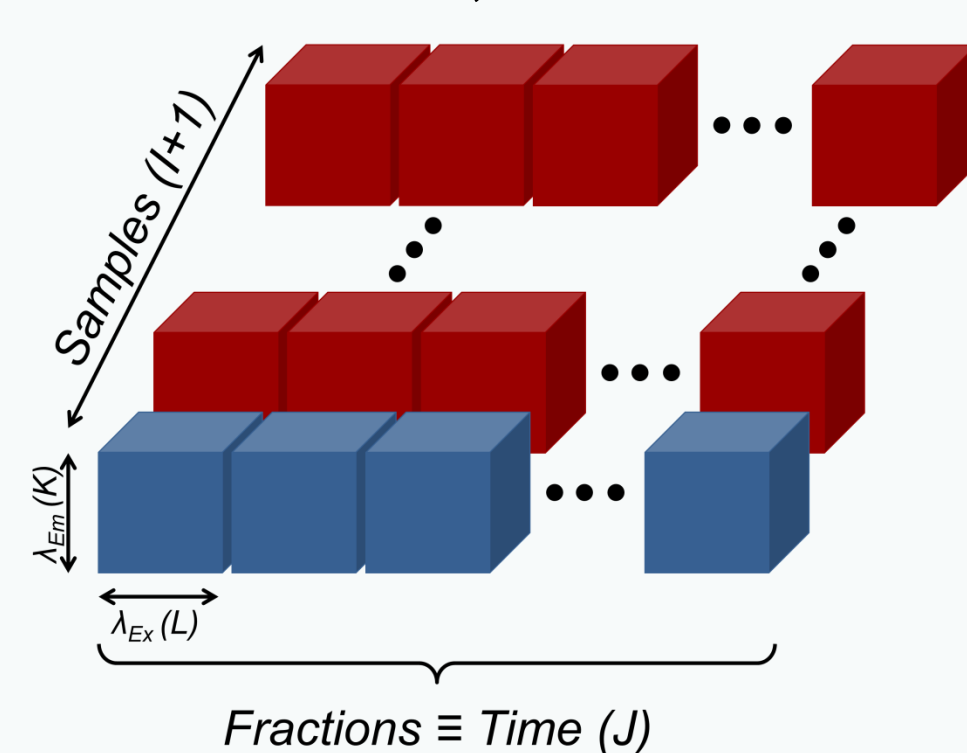
(J×K×L): 25×17×25

Data pre-processing:

EEM* smoothing

Data processing:

PARAFAC, APARAFAC, MCR-ALS, U-PLS/RTL



*EEM= Excitation-Emission Matrix

Analysis time per sample

40 min

Matrix size

(J×K×L): 10×121×25

Data pre-processing:

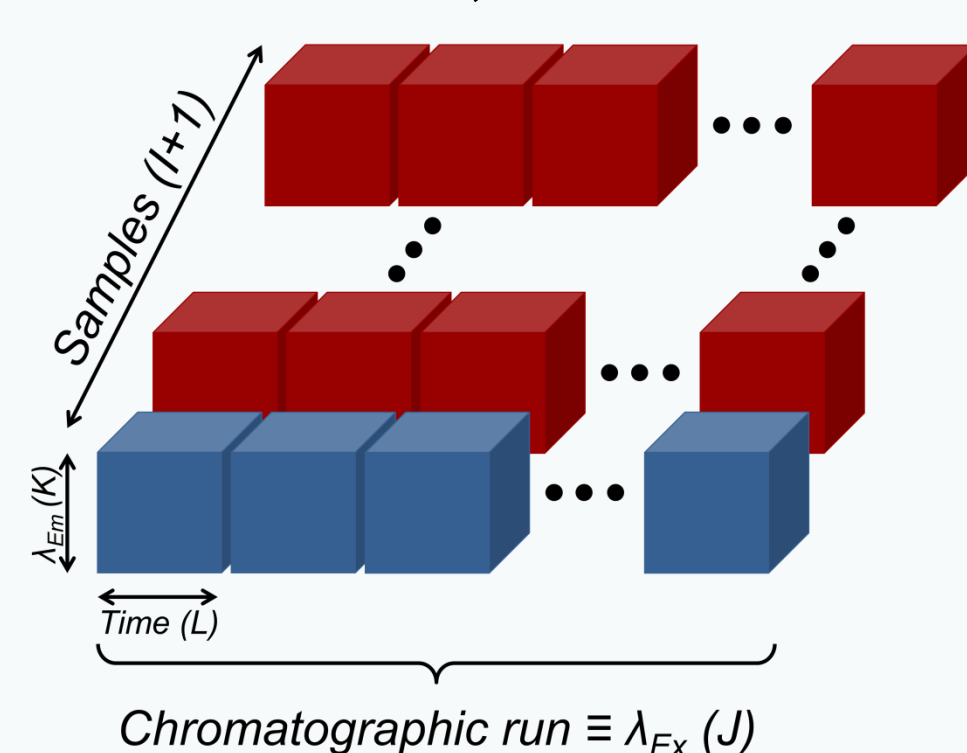
TEM* smoothing

Peak alignment

TEM baseline correction

Data processing:

PARAFAC, APARAFAC, MCR-ALS, U-PLS/RTL



*TEM= Time-Emission Matrix

Analysis time per sample

(HPLC + FD)= 7 min

Matrix size

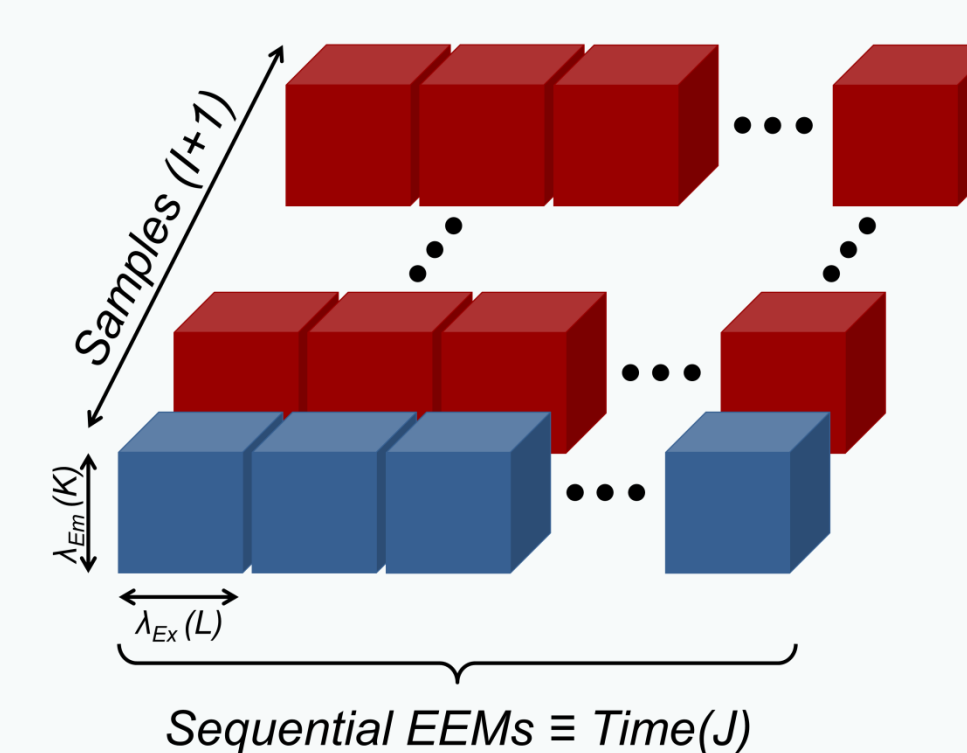
(J×K×L): 15×28×15

Data pre-processing:

TEM smoothing

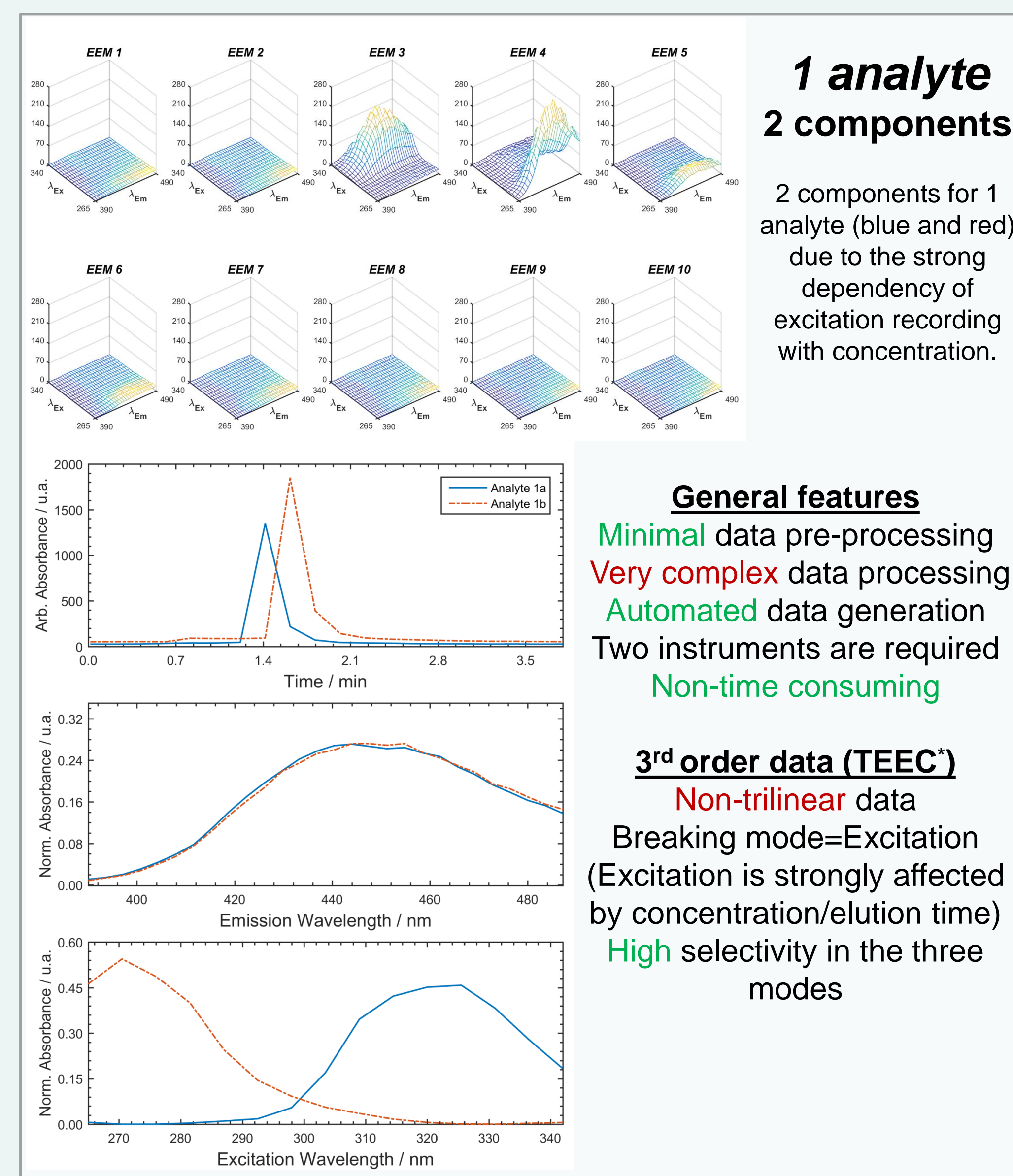
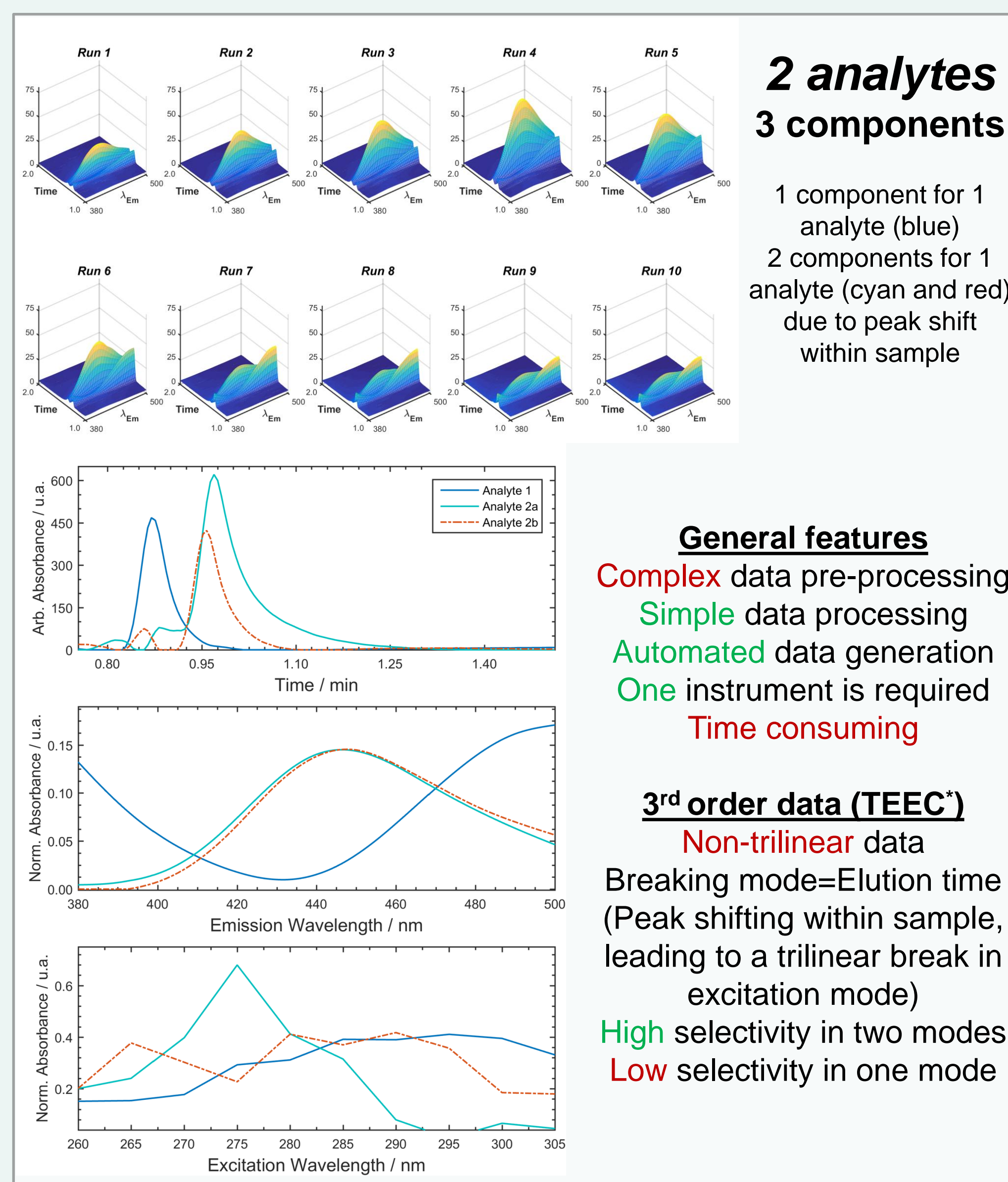
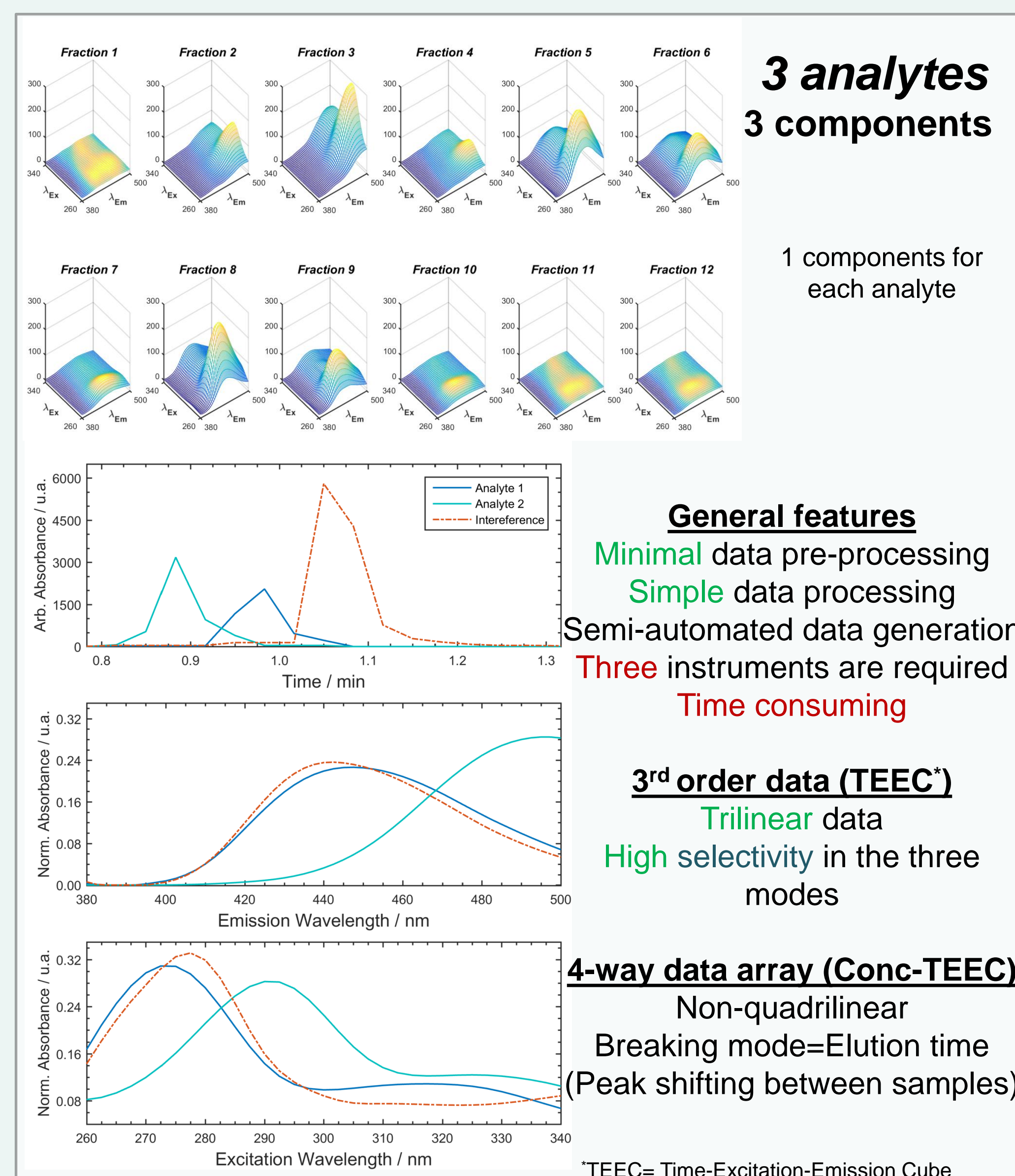
Data processing:

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RESULTS and DISCUSSION

In order to demonstrate the complexity of the system, the data pre-processing just included smoothing. No peak alignment or baseline correction was performed. The figures show PARAFAC results.



CONCLUSIONS

The three techniques allow to obtain third-order data in a simple way without using sophisticated instrumentation. **Method 1** becomes the selected one to generate this kind of data despite of the instrumentation requirement and the time consumption. The obtained data is simple and well-known algorithm are used for its modelling. **This methodology could be improved by using a fast-scanning spectrofluorometer.** On the other hand, just a HPLC with fast-scanning fluorescence detector is required for **Method 2**. However, due to its high time consumption, it is not appropriate for long chromatographic runs and analysis of unstable compounds. Furthermore, complex data pre-processing could be demanded prior to data modelling. Last but not least, the **Method 3** shows the most promising methodology for 3rd-order data generation, considering its equipment simplicity and fast data acquisition. Nevertheless, **it is strongly necessary to develop an algorithm that allows to model this data, taking into account its particular characteristics.**