



A focus for analytical chemistry in Europe



WG. 1.3. Recent instrumental developments in targeted and non-targeted analysis.

Chair
Ivo Leito

20.-21. May 2019, Tartu, Estonia

Validation of a non-target method: a practical approach



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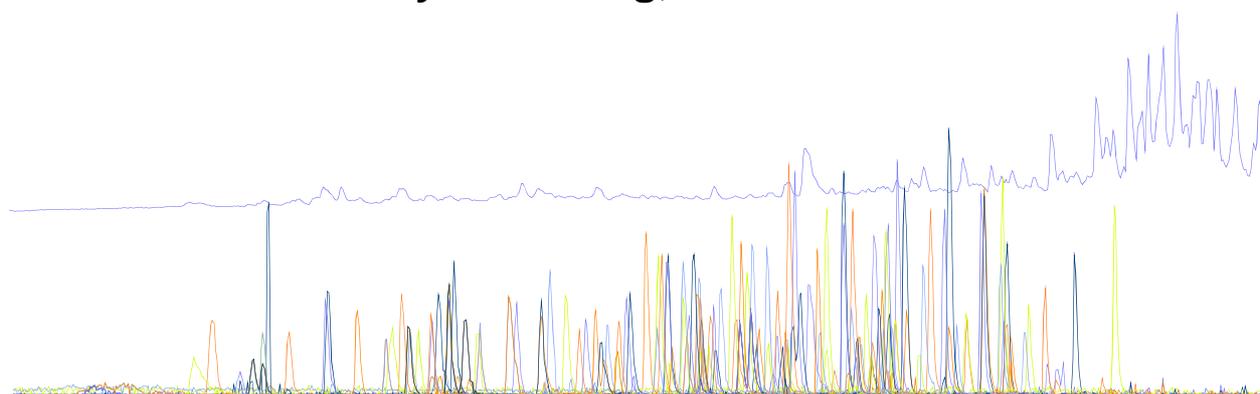
- Introduction
 - 31 participants, close to 20 countries
 - From universities, institutes, authorities, labs, industry
- How was workshop arranged: any deviation into groups? **No**
how many questions for discussion? **Mostly from chair**

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A typical picture:

Food Safety Screening, QuEChers Extract



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Scheme: Sven Vedder, Shimadzu

Non-target screening with high resolution mass spectrometry: Critical review using a collaborative trial on water analysis

Emma L. Schymanski^{1*}, Heinz P. Singer¹, Jaroslav Slobodnik², Ildiko M. Ipolyi², Peter Oswald², Martin Krauss³, Tobias Schulze³, Peter Haglund⁴, Thomas Letzel⁵, Sylvia Grosse⁵, Nikolaos S. Thomaidis⁶, Anna Bletsou⁶, Christian Zwiener⁷, María Ibáñez⁸, Tania Portolés⁸, Ronald de Boer⁹, Malcolm J. Reid¹⁰, Matthias Onghena¹¹, Uwe Kunkel¹², Wolfgang Schulz¹³, Amélie Guillon¹⁴, Naïke Noyon¹⁴, Gaëla Leroy¹⁵, Philippe Bados¹⁶, Sara Bogialli¹⁷, Draženka Stipaničev¹⁸, Paweł Rostkowski¹⁹, Juliane Hollender^{1,20*}.

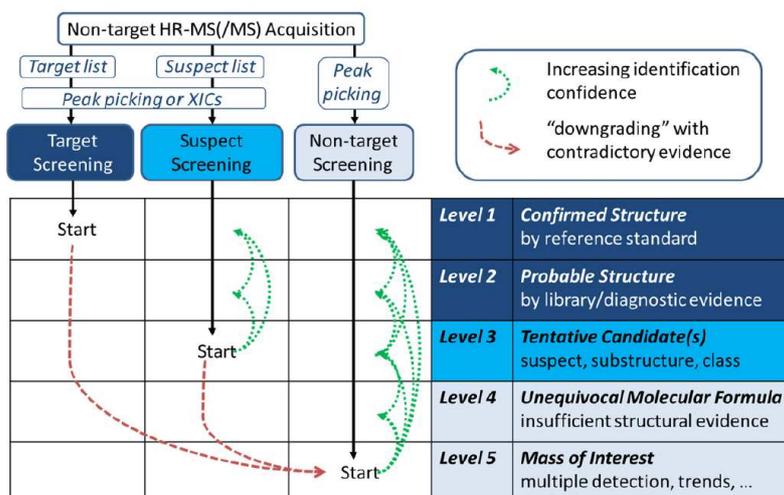


Figure 1: Matrix of identification approach versus identification confidence.

[Analytical and Bioanalytical Chemistry](#)

August 2015, Volume 407, Issue 21, pp 6237–6255

Scheme: Sahun Bilsborough, Agilent

Workflow



- **Reproducibility, robustness**
- **Sample preparation**

- **As routine as possible**
 - Suspect list
 - That defines the method
 - E.g. 300 pesticides on the list

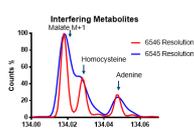
- **Identification**
 - RT, m/z accuracy, isotope ratio accuracy, MS/MS spectral match

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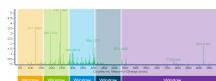
Possible workflow:



High resolving power even at high acquisition speeds



QRAI workflow to acquire MS and MS/MS spectra in a single run with increased separation and LOQ



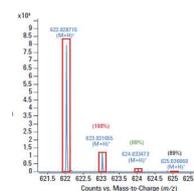
Software workflows to maximise productivity in your laboratory

4 confirmation criteria for suspect screen

Wide in spectrum dynamic range to enable ID of high and low abundance compounds in a single analysis



High quality isotopic fidelity to increase confidence in identifications



Scheme: Sahun Bilborough, Agilent

Instrumental



- **When targeted at low LoD: QqQ**
- **When non-targeted, complex samples: Q-TOF, triple-TOF** seem most practical
 - **HRAM, Resolution** $n \times 10\,000$
 - **Wider dynamic range** > 4 orders
 - Form one run vastly different
 - **Accurate isotopic patterns**
 - Ca 95% false positive scan be eliminated
 - **Advanced acquisition**
- Ion sources
- LC

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Acquisition and software



- **Software is BIG**
- Handling **large number of components**
 - In LC mode
 - Algorithms for detecting peaks
 - Advanced data acquisition strategies (SWATH, ...)
 - Open software, open platforms
 - Identification
 - RT, m/z accuracy, isotope ratio accuracy, MS/MS spectral match
- **Databases**
 - Curation (removal of imperfections)
- **Statistical analysis**

Nothing is perfect!

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