TUTORIAL ON 'DATA QUALITY AND SMART DATA HANDLING IN FOOD ANALYSIS'

L105

DATA PROCESSING AND IDENTIFICATION OF SMALL MOLECULES IN LC-MS-BASED NON-TARGETED ANALYSIS WORKFLOWS

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Presentation will focus on:

- Overview of non-targeted workflows in food analysis (Workflows and terminology)
- LC-MS platforms and approaches to non-targeted data acquisition
- Requirements and assurance of HR-MS data quality (Mass locking, re-calibration of mass spectra, RT normalization, internal standards,...)
- Export of data for chemometric handling
- Data mining, pre-processing and analysis (Tools, methods, (un)supervised pattern recognition,...)
- Identification of small molecules: approaches and tools (Elemental formula generation, mass spectral libraries and chemical databases, prediction and
 - interpretation of mass spectra,...)
- Examples
- General recommendations

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CRITICAL REVIEW, EXPERIENCES AND OUTLOOK WITH RESPECT TO METABOLOMICS DATA HANDLING OPTIONS

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Presentation will focus on key parameters in ensuring quality of LC-HRMS metabolomics data and models

- Overview of LC-MS based metabolomics workflow (Identification of critical steps)
- Quality of the samples (Key parameters in designing the experiment)
- Analytical Quality (Expectations and requirements in both sample preparation and fingerprinting steps)
- Data Analysis Quality (Tools for validating the models / markers)
- Illustrations and General recommendations to ensure robustness of the whole workflow