



LIVE WEBINAR EXPAND YOUR METABOLOMICS HORIZONS

Live Webinar hosted by Separation Science

WEBINAR DETAILS

Date: 14 September, 2016

Start Time: This event will broadcast live at two separate times - select the most convenient time:

9 a.m. EDT / 2 p.m. BST / 3 p.m. CEST

12 p.m. PDT / 3 p.m. EDT

Metabolomics is a powerful technique for understanding biological systems by measuring the abundance of metabolites. The field is rapidly evolving as new approaches are developed to the analytical and biological challenges researchers face. Three workflow improvements will be discussed in this presentation:

- Advances in batch feature extraction software (Profinder) for LC/MS & GC/MS data
- A robust and reproducible LC/QQQ method and acquisition database for the targeted measurement of >215 central carbon metabolites
- Software solution for qualitative flux analysis (stable isotope tracing) by LC/Q-TOF

What Will You Learn?

New approaches to understanding the biological context of diseases and the ability to map quantitative results onto biological pathways. How to extend the power of your metabolomics research by studying the metabolic network in motion and interpret results for communication with colleagues using pathway images and videos of qualitative fluxes.

Register now

[www.agilent.com/chem/
MassSpec_MetaWebinar](http://www.agilent.com/chem/MassSpec_MetaWebinar)

For Research Use Only. Not for use in diagnostic procedures. This information is subject to change without notice.

© Agilent Technologies, Inc. 2016
Printed in USA, August 16, 2016
5991-7306EN

