

The background of the top half of the slide is a vibrant blue and green abstract image. It features glowing molecular structures, circuit-like patterns, and a sense of dynamic energy. A large, light-blue circle with a thin black border is positioned on the right side, containing the event title.

## Innovate with Mass Spectrometry: Lunch & Learn Series

Wednesday, March 20th, 2019

9:00 AM-2:00 PM Eastern

Location: UCONN-Storrs Chemistry Building room A304

### Agenda

**9:00 AM** The Basics of LC Mass Spectrometry | Dr. Adam Graichen, University of CT

**9:30 AM** Advances in reverse phase chromatography |  
Dani Xing, Phenomenex

**10:00 AM** Coffee Break

**10:30 AM** Lipidomics and metabolomics with HILIC chromatography | Bob Proos, SCIEX

**11:00 AM** Simultaneous Targeted and non-Targeted Analysis in Citrus with X500R QTOF | Craig Butt, SCIEX

**11:30 AM** Improving Drug Metabolite Identification Studies by Optimizing Acquisition, Ion mobility and Processing Parameters | Rick Schneider, Pfizer (retired)

**12:00 PM** Lunch

**1:00 PM** Industrialized Proteomics using the TripleTOF® 6600 and QTRAP® 6500+ | Mahbod Hajivandi, SCIEX

**1:30 PM** Polly: A Comprehensive Analysis and Workflow Platform for the Metabolomics Lab |  
Jim Martin Elucidata

**2:00 PM** Close

Register today at <https://info.sciex.com/lcms2019>

\* The opinion of the invited speakers are theirs and do not necessarily reflect the views of SCIEX.



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