



Wednesday, March 26, 2025 · 11:30 a.m.-1:00 p.m.

Benjamin P. Boussert Conference Room | LSU
100 Chemistry & Materials Building

Join us and learn how targeted quantitative LC-MS(/MS) workflows can be impactful for your research. We will highlight some of the newest technology to help you be even more successful. Oh, and lunch is on us!

Discussion Topics:

From Discovery to Targeted Analysis

Translating qualitative studies to targeted panels for larger biological cohort studies

Targeted Analysis for Lipids ,Fatty Acids, and metabolites

Identification and quantification of lipid classes and endogenous and novel metabolites

Small Molecule Quantitation for Specific Molecule classes including Analgesics

Accurate quantitation for targeted small molecule including pharmaceuticals

Protein and peptide characterization and quantitation

Characterization of proteins including post transitional modifications and protein-protein interactions and peptides identification and quant

Meet Your Team

Rachel Kidwell, LCMS,
Rachel.Kidwell@sciex.com
m 903.216.0426
www.sciex.com

Christa Feasley, PhD
Advanced Workflow Specialist
christa.feasley@sciex.com
www.sciex.com

Please register with the QR
Code for lunch!





Get to know your SCIEX speaker!



Christa Feasley, PhD

Advanced Workflow Specialist supporting the South-Central and Southeast sales districts, SCIEX

Christa current focus is on SCIEX high-resolution mass spectrometry platforms particularly driving regional growth of Life Science Research (LSR), Pharma, and Biopharma business.

Christa has multiple mass spec roles prior to joining SCIEX. She earned her Ph.D. in Chemistry from Purdue University, then established a core laboratory for glycoconjugate characterization at University of Oklahoma Health Sciences Center. From there, she joined Thermo as an applications scientist focused on both pre- and post-sales customer support with focus in the biopharma market. Most recently, Christa was a senior analytical development scientist at Cytovance Biologics, a CDMO in Oklahoma City, where she built a new mass spec services unit to support protein characterization.