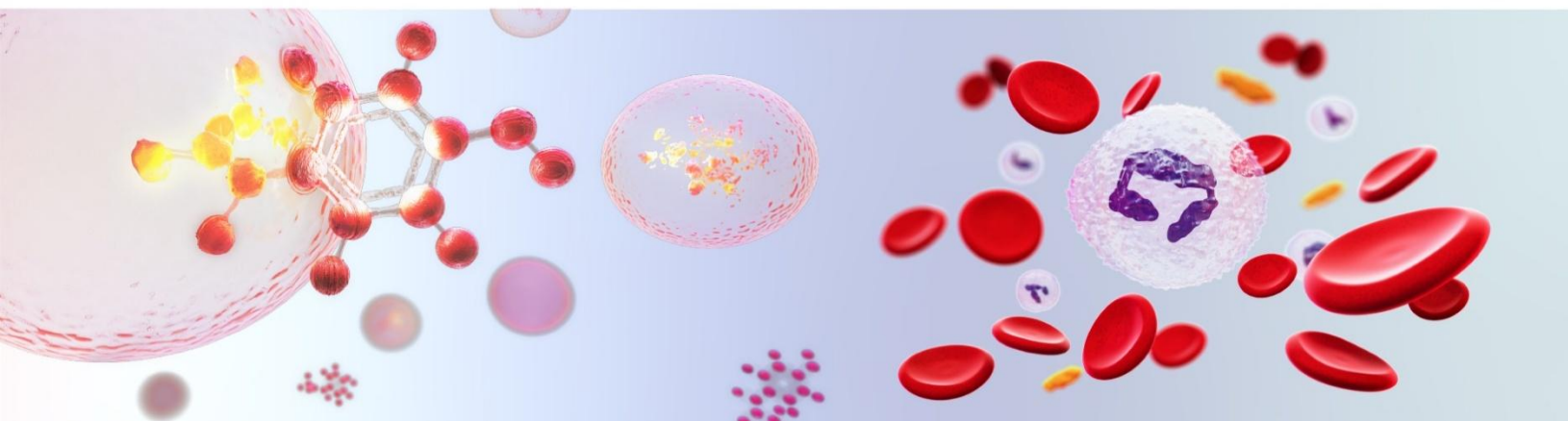


Discovery continuum seminar

Tuesday, August 18, 2026 | 8:30 a.m. – 5:00 p.m.

Green Acre Campus Point | 10300 Campus Point Dr | San Diego, CA 92121



Join SCIEX and industry leaders for an in-depth seminar designed to address the most pressing challenges in drug discovery and development. This event covers the full spectrum of analytical solutions driving progress in pharma and biopharma.

Agenda:

- 8:30 a.m. Registration and breakfast
- 9:00 a.m. Welcome and introductions – Josh Plath, SCIEX
- 9:10 a.m. Innovations in mass spectrometry and software: Unveiling the latest SCIEX advancements – Christa Feasley, SCIEX
- 9:40 a.m. Ultra-high throughput MS analysis in oligonucleotide manufacturing – Davis Hickey, Illumina
- 10:10 a.m. Ultra-high-throughput metabolic stability assay powered by acoustic ejection mass spectrometry with sample-specific MS/MS methods – Craig Gutierrez, Iambic
- 10:40 a.m. Coffee break
- 11:00 a.m. AI Quantitation software delivers powerful automated precision – Elliott Jones, SCIEX
- 11:30 a.m. LCMS strategies for successful pharmacokinetic assessment of next-generation ADCs – Dave Beyerlein, BioAgilytix
- 12:00 p.m. The power of insight: How polar integration transforms data into decisions – Shendi Lu, IDBS
- 12:30 p.m. Lunch and networking
- 1:15 p.m. Navigating mass spectrometry challenges in the drug discovery continuum – A CRO perspective – Shantanu Roychowdhury, Eurofins Discovery Services, North America, LLC
- 1:45 p.m. In-depth characterization of trastuzumab and its conjugated ADC with multi-level icIEF-UV/MS analysis – Jingwen Ding, SCIEX
- 2:15 p.m. Optimization of capillary electrophoresis method for purity analysis of large synthetic self-replicating RNA – Jacqueline Posy, Replicate Bioscience, Inc.
- 2:45 p.m. Characterization of peptide therapeutics and conjugated protein modalities – Vikram Shenoy, Phenomenex
- 3:15 p.m. Closing remarks
- 3:30 p.m. Happy hour refreshments
- 5:00 p.m. Event adjourns



Register today: <https://sciex.li/discovery-continuum-SDCA>

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