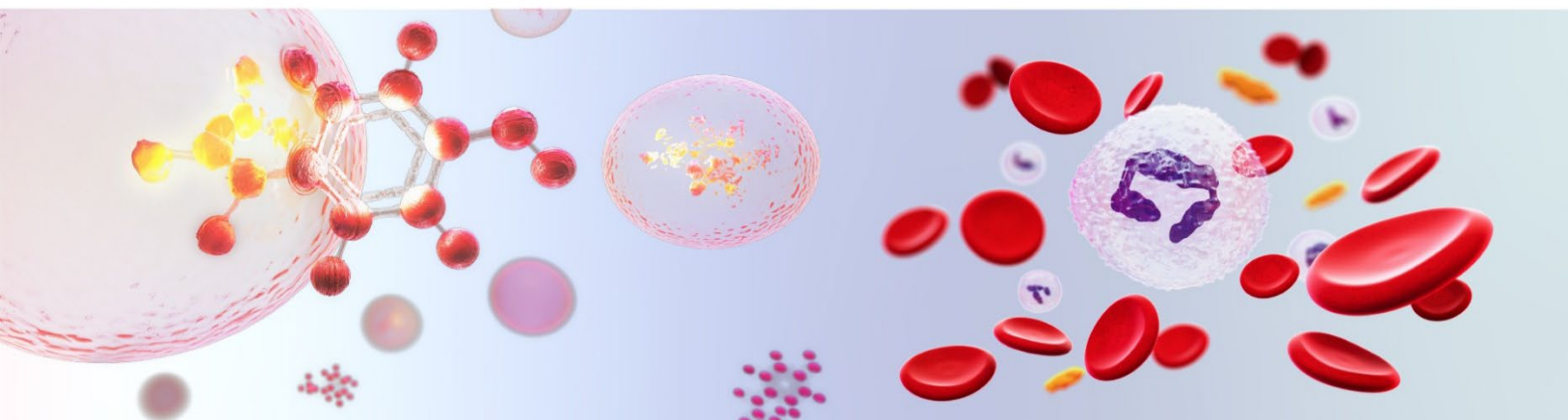


# Discovery continuum seminar

Tuesday, June 16, 2026 | 8:30 a.m. - 5:00 p.m.  
MIT Museum | 314 Main St. | Cambridge, MA 02142



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 – James Dougherty and Josh Plath, SCIEX
- 9:10 a.m. Advances in AI-powered quantitative analysis – Jose Castro-Perez, SCIEX
- 9:40 a.m. Automated HT-ADME: SCIEX 7600 system for AI driven drug discovery – Li Li, Cyprotex
- 10:10 a.m. Leveraging the Echo® MS+ with ZenoTOF to build the lab of tomorrow – Hilary Ranson, Takeda
- 10:40 a.m. Coffee break
- 11:00 a.m. Multi-summed DPPT–HRMS/MS for ultra-sensitive quantification of diverse siRNA conjugates in plasma and tissues – Dan Kim, City Therapeutics
- 11:30 a.m. The analysis of immunoglobulins glycosylation patterns in plasma samples using EAD on the ZenoTOF 7600+ system – Bogdan Budnik, Wyss Institute
- 12:00 p.m. SysQuan: A novel and affordable approach for absolute quantitation of the human and mouse tissue specific proteome – Christoph Borchers, Mcgill University
- 12:30 p.m. Lunch and networking
- 1:15 p.m. Investigating mispairing heterogeneity in bispecific antibodies by ultra-high resolution icIEF-UV/MS – Tian Xu, Lilly
- 1:45 p.m. Onboarding the Biophase 8800 system for use at KBI Biopharma – David Abraham, KBI Biopharma
- 2:15 p.m. A brave new world: Walking on the wild side [of software] – Cindy Sanderson, SCIEX
- 2:45 p.m. Dead code walking: Why unsupported software is a silent data integrity risk – Milind Nadgouda, RiverArk Limited
- 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-MA>

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