



4th Cambridge Proteomics and Mass Spectrometry Meeting
(11am for AGM) 12-5pm Thursday 23rd April 2026

Cambridge Building, Babraham Research Campus, Cambridge

Please note: registration is essential, and we have no space for posters this time.

—10.30 am— Refreshments, networking, setup—

11.00am Our first CamsPaMS AGM - all attendees, feedback, and ideas for the future of the meeting welcome!

—12.00 pm— Lunch, networking—

Session 1—Chair: Richard Kay

12.45 pm Welcome and meeting introduction

12.50 pm ZenoTOF 8600 - redefining sensitivity in accurate mass spectrometry (Sponsored) **Tom Ruane, SCIEX**

1.00 pm Unmasking the perfect switch with msHDX-MS **Katie Gibson, AZ**

1.20 pm Enhanced Spatial Resolution of the Toxoplasma gondii Proteome by HyperLOPIT-XL **Alex Lester, Uni of Cambridge**

—1.40 - 2.15 pm— Refreshments, networking—

Session 2—Chair: Ashley Phillips

2.15 pm Olink (Sponsored)

2.20 pm Overcoming biotherapeutic-induced interference in a multiplexed immunoaffinity LC-MS assay for the quantitation of PAD2 and PAD4 in human serum **Yuki Long, Resoliant**

2.40 pm GSK dia-PASEF Enables Rapid Profiling of the Human Secretome for Deeper Insights into Cellular Dynamics and Inflammatory Mechanisms **Chloe Tayler, GSK**

3.00 pm Old drugs and new challenges - a review of nitazenes in the UK **Simon Hudson, LGC Ltd**

—3.20 - 3.50 pm— Refreshments, networking—

Session 3—Chair: Marcus Ladds, Jo Kirkpatrick

3.50 pm **Roundtable discussion: Single Cell Mass Spectrometry Technologies**

Panel members: Mel Bailey (King's College London), Alex Van Der Walt (Tecan), Ian Fahy (Beckman Coulter), Georgina Charlton (GSK)

4.45 pm Wrap up and quiz

—5.00 pm onwards— Pizza, drinks, networking in the bar downstairs—

—7.00 pm onwards— Further networking at the George pub, Babraham—