

EPIC-XS

Programme: Thursday 28 October 2021, 15:40 - 18:00
(14:40-17:00BST, 10:40-13:00 EST, 07:40-10:00 PST)

Welcome	
15:40 -15:45	Welcome from the Chairs, Eduard Sabidó , Centre for Genomic Regulation, Spain Emma Lundberg , Science for Life Laboratory, Stockholm, Sweden

Session 1	Gene regulation and protein turnover Chair: Eduard Sabidó
15.45 - 16.05	RNA binding proteins in melanoma progression Fatima Gebauer , Centre for Genomic Regulation, Spain
16.05 - 16.25	Tracking down the elusive interactors of Polycomb proteins by highly sensitive tissue proteomics in Drosophila Giacomo Cavalli , Institute of Human Genetics, France
16.25 – 16.45	Quantifying and understanding protein turnover control by DIA-MS Yansheng Liu , Yale University School of Medicine, USA
16.45 – 17.00	Q&A with the 2 speakers, moderated by Eduard Sabidó

17.00 – 17.10 Coffee break



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Session 2	Immunopeptidomics and sequencing technologies Chair: Emma Lundberg
17:10 - 17.30	Deciphering the HLA-peptidomics landscape in melanoma. Yardena Samuels , The Weizmann Institute of Science, Israel
17.30 - 17.50	Single-molecule protein sequencing Chirlmin Joo , Delft University of Technology, The Netherlands
17.50 – 18.00	Q&A with the speakers, moderated by Emma Lundberg