

Programme: Thursday 08 Sept. 2022, 15.00 – 17.30CET
 (14:00-18:05 BST, 09:00-11:30 EST, 06:00-08:30 PST)

Welcome	
15:00 -15:10	Welcome from the Chair, Tiziana Bonaldi , EPIC-XS – IEO, Italy
Session 1	Chair: Tiziana Bonaldi
15.10 - 15.25	Characterizing ADP-ribosylation in health and disease using quantitative mass spectrometry - Michael Lund Nielsen , CPR- Copenhagen
15.25 - 15.40	Plant HCN Regulates Cellular Processes through a new Posttranslational Modification of Proteins: S-cyanylation - Irene Garcia CSIC, Sevilla
15:40 – 15:55	About Protein Histidine Phosphorylation (& using IMAC for XL-MS) - Albert Heck , Utrecht University
15.55 – 16.10	Q&A with the 3 speakers, moderated by Kris Gevaert
16.10 - 16.20 Coffee break	
Session 2	Chair, Kris Gevaert
16.20 – 16.35	Role of arginine-methylation in adaptive response of cancer cells to cisplatin - Marianna Maniaci , IEO, Milan
16.35 - 16.50	Unbiased exploration of the proteome-wide modification landscape - Lennart Martens , VIB-Ghent
16.50 – 17.25	Keynote Speaker – Neil Kelleher , Northwestern University To dial in on a nucleosome code: take it from the top down using native, Nuc-MS
17.25 – 17.30	Q&A with the 3 speakers, moderated by Kris Gevaert