Day1, Tue 13.09.22					Day 2,Wed 14.09.22					Day 3, Thu 15.09.22				
time	type	name	affiliation	title	time	type	name	affiliation	title	time	type	name	affiliation	title
9:00	intro	Klaus Kratochwill	President of APMA, Head of Core Facility Proteomics at Medical	Welcome words	9:00	D2_invited 1	Fan Liu	Head of Masspec, FMP Leibniz, Berlin, Germany	Developing proteome-wide cross-linking mass spectrometry and its application in biology	9:00	D2_inviteD 2	Alexander Makarov	Director Global Research LSMS at Thermofisher Scientific, Bremen, Germany	Orbitrap mass spectrometry for single-cell proteomics
9:15					9:15					9:15				
9:30	D1_invited 1	Josef Penninger	Institute, University of British Columbia, Vancouver, Canada	Keynote Lecture, TBA	9:30	D2_invited 2	Anjali Seth	Head of Single Cell Proteomics at Cellenion,	TBA	9:30				
9:45					9:45					9:45	D2_invited 2	TBA	HP	TBA
10:00					10:00	D2_short_talk_1				10:00				
10:15	D1_short_talk1				10:15					10:15				
10:30					10:30	coffee break				10:30	D2_short_talk1			
10:45	coffee break				10:45					10:45				
11:00					11:00	D2_invited 3	Erwin Schoof	Technical University of Denmark	TBA	11:00	coffee break			
11:15	D1_invited 2	Roman Zubarev	Professor of medicinal proteomics in the Department of Medical Biochemistry and Biophysics,	Introduction into mass spectrometry	11:15					11:15				
11:30					11:30					11:30	D1_invited 3	Ákos Végvári	Karolinska Institutet, Stockholm, Sweden	Data acquisition strategies for improved protege profiling in single cell proteomics
11:45					11:45	D2_invited 4	Bogdan Budnik	Wyss Institute at Harvard University, Boston, MA, US	Single-cell Multi-omics and PTM-omics are the next two challenges that need to be resolved in our march towards a comprehensive digital Pathomics	11:45				
12:00	D1_short_talk2				12:00					12:00				
12:15					12:15					12:15	D1_short_talk2			
12:30	lunch break				12:30	lunch break				12:30				
12:45					12:45					12:45	lunch break			
13:00					13:00					13:00				
13:15					13:15					13:15				
13:30	D1_invited 4	Robert Ahrends	Institute of Analytical Chemistry, University of Vienna, Austria	TBA	13:30	D2_invited 5	Benjamin Furtwängler	University of Copenhagen, Copenhagen	Real-time search assisted single-cell proteomics applied on human hematopoiesis	13:30				
13:45					13:45					13:45	D1_invited 4	TBA	Bruker	TBA
14:00					14:00					14:00				
14:15	D1_invited 5	Ruth Birner Grünnberger	Institute of Chemical Technologies and Analytics, TU Vienna, Austria	Lipid hydolysis in cancer	14:15	D2_invited 6	Manuel Matzinger	IMP, Vienna	Establishment of reproducible, sensitive and automated one-pot workflows for high throughput single cell proteomics.	14:15				
14:30					14:30					14:30	D1_short_talk3			
14:45					14:45					14:45				
15:00	D1_short_talk3				15:00	D2_short_talk2				15:00	D1_short_talk4			
15:15					15:15					15:15				
15:30	coffee break				15:30	coffee break				15:30	coffee break			
15:45					15:45					15:45				
16:00	D1_invited 6	Roman Zubarev	Professor of medicinal proteomics in the Department of Medical Biochemistry and Biophysics,	Single Cell Chemical Proteomics	16:00	D2_invited 6	Chris Rose	Genentech	Intelligent data acquisition strategies for single cell proteomics	16:00	D1_invited 6	Ryan T. Kelly	Brigham Young University, Provo, UT, US	Comparison of data acquisition strategies for label-free single-cell proteomics
16:15					16:15					16:15				
16:30					16:30					16:30				
16:45	D1_short_talk4				16:45	D2_invited 7	Ying Zhu	Pacific Northwest National Laboratory, Richland, WA	Advancing sample preparation and MS data acquisition for single-cell and spatial multiomics	16:45	closing	Karl Mechtler	IMP, Vienna	closing remarks
17:00					17:00					17:00				
17:15	D1_short_talk5				17:15									
17:30					17:30	D2_short_talk3								
					17:45									
						Conference Dinner								