## PROGRAM

## **MONDAY, 20 MAY 2019**

08:00-09:00	Registration
09:00-09:15	Introduction by organizers
SESSION 1	UPS, the Ubiquitin Proteasome System
09:15-10:00	50 Years of Degradation: Looking Back and Looking Forward ALFRED GOLDBERG, Harvard Medical School, Boston, USA
10:00-10:15	Ubp6 and Rpn11 broad and unique specificities place them as master regulators of protasomal substrates processing
T01	DARIA RIABOV BASSAT, Weizmann Institute of Science, Israel
10:15-10:30	Regulation of histone degradation in response to DNA damage by the Ubiquitin-proteasome system
T02	KIRAN CHALLA, FMI Basel, Switzerland
10:30-11:00	Coffee break
11:00-11:30	Mechanistic insights into pupylation, a bacterial Ub-like modification pathway EILIKA WEBER-BAN, ETH Zurich, Switzerland
11:30-11:45 <mark>T03</mark>	Cancer Cell Vulnerability to SMARCA2/4 Degradation by PROTACs MANFRED KOEGL, Boehringer Ingelheim, Vienna, Austria
11:45-12:15	Flash talks 1 – even numbers
12:15-13:45	Lunch with poster session 1 (even numbers)
12:15-13:45 13:45-14:00	Lunch with poster session 1 (even numbers) Group photo
13:45-14:00	Group photo
13:45-14:00 SESSION 2	Group photo Protein Quality Control Quality control at the ribosome during translation
<b>13:45-14:00</b> SESSION 2 14:00-14:30 14:30-14:45	Group photo Protein Quality Control Quality control at the ribosome during translation RAMANUJAN HEGDE, MRC Laboratory of Molecular Biology, Cambridge, UK Protein Homeostasis at the Golgi Apparatus DORIS HELLERSCHMIED, Yale University, University Duisburg/Essen,
13:45-14:00           SESSION 2           14:00-14:30           14:30-14:45           T04	Group photo Protein Quality Control Quality control at the ribosome during translation RAMANUJAN HEGDE, MRC Laboratory of Molecular Biology, Cambridge, UK Protein Homeostasis at the Golgi Apparatus DORIS HELLERSCHMIED, Yale University, University Duisburg/Essen, Germany Unique mechanisms for maintaining protein-folding homeostasis in the endoplasmic reticulum

SESSION 3	Molecular Functions and Tools
16:00-16:30	LUBAC and linear ubiquitin chains: novel tools to study immune signaling KATRIN RITTINGER, The Francis Crick Institute, London, UK
16:30-16:45	A tri-ionic anchor mechanism drives Ube2N-specific recruitment and K63-chain ubiquitination in TRIM ligases
<b>T06</b>	LEO KISS, MRC Laboratory of Molecular Biology, Cambridge, UK
16:45-17:30	Flash talks 2 – odd numbers
17:30-20:00	Reception with poster session 2 (odd numbers)
20:30	Conference dinner

## **TUESDAY, 21 MAY 2019**

SESSION 4	Ubiquitin and Friends
09:00-09:45	Unconventional Serine Ubiquitination IVAN DIKIC, Goethe University, Frankfurt, Germany
09:45-10:00 <b>T07</b>	A high-fidelity multiplex CRISPR/Cas library for functional interrogations within the autophagy network <b>MANUEL KAULICH</b> , Goethe University Frankfurt, Germany
10:00-10:30	Distinct functions of ATG16L1 isoforms in membrane binding and LC3B lipidation in autophagy-related processes ANNE SIMONSEN, Institute of Basic Medical Sciences, Oslo, Norway
10:30-11:00	Coffee break
SESSION 5	Emerging Ubiquitin Cell Biology
SESSION 5 11:00-11:30	Emerging Ubiquitin Cell Biology Ubiquitination of the E3 ligase HOIP regulates immune signaling and cell death FUMIYO IKEDA, Institute of Molecular Biotechnology, Vienna, Austria
	Ubiquitination of the E3 ligase HOIP regulates immune signaling and cell death

12:15 Closing remarks