Science Talks Campus Solna SciLifeLab research symposium 2020



Session I - 26 august 2020 Moderator: Per Ljungdahl

09:00	Opening note Per Ljungdahl, Campus Solna Director
09:10	GPSeq reveals radial genome organization at high resolution Magda Bienko, Karolinska Institutet
09:30	Illuminating the origins of DNA copy number variation by single-cell sequencing Nicola Crosetto, Karolinska Institutet
09:50	Understanding the importance of dynamics in efficient plastic degradation by designer enzymes Per-Olof Syrén, KTH
10:10	Intestinal estrogen receptor beta suppresses colon in lammation and tumorigenesis in both sexes

Amena Archer, KTH

10:30 Break out rooms - meet the speakers

10:45 Coffee Break

Session II

Moderator: Simon Elsässer

11:00	Human systems immunology Petter Brodin, Karolinska Institutet
11:20	Drug discoveries and public-private partnerships in translational microbiome research Lars Engstrand, Karolinska Institutet
11:40	Single cell proteome heterogeneity of human cells Emma Lundberg, KTH
12:00	Lung cell heterogeneity and its impact on tissue plasticity Christos Samakovlis, Stockholm University
12:20	Break out rooms - meet the speakers
12:35	Short break

Session II

Moderators: Christian Sommerauer & Aswathy Kallazhi

12:40	GPSeq reveals the radial landscape of chromatin and genomic alterations in the human cell nucleus Gabriele Girelli, Karolinska Institutet
12:50	Deciphering the roles of sex hormones in the human immune system Camila Consiglio, Karolinska Institutet
13:00	Characterization of genetic regions at high risk for differential population response across chemical classes Marissa Kosnik, Stockholm University
13:10	Break out rooms - meet the speakers
13:25	End of session

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Session IV- 27 august 2020 Moderator: Alexey Amunts

09:00 Longitudinal analysis of molecular and digital health profiles during lifestyle changes

Francesco Marabita, Karolinska Institutet

09:20 The use of systems biology in treatment of liver diseases

Adil Mardinoglu, KTH

09:40 Views and clues about the circulating proteome

Jochen Schwenk, KTH

10:00 Exploring the human proteins to understand biology and disease and to develop therapeutic

drugs

Mathias Uhlén, KTH

10:20 Break out rooms - meet the speakers

10:35 Coffee Break

Session V

Moderator: Kristina Jonas

10:50 Aquatic microbial diversity, functions and interactions

Sarahi Garcia, Stockholm University

11:10 A ribosome pause-site regulates the translation of the bacterial DNA replication initiator

DnaA in response to nutrient starvationMichele Felletti, Stockholm University

11:30 Investigating ribosome dynamics by 5'P mRNA degradome sequencing

Vicent Pelechano, Karolinska Institutet

11:50 Exploring the small proteome

Simon Elsässer, Karolinska Institutet

12:10 Break out rooms - meet the speakers

12:25 Short break

Session VI

Moderators: Dörte Schlesinger & Eva Brinkman

12:30 Site specific incorporation of two non-canonical amino acids for two-colour bioothorgonal

labelling and chemical-controlledBirthe Meineke, Karolinska Institutet

12:40 Large-scale identification of RBP-RNA interactions by RAP-seq refines essentials of post-

transcriptional gene regulationIonut Atanasoai, Karolinska Institutet

12:50 Tandem talk: A LAMP assay for detection of SARS-CoV-2 using in-house produced enzymes

Yerma Pareja-Sánchez, Karolinska Institutet & Rebecca J Howard, Stockholm University

13:00 Break out rooms - meet the speakers

13:15 End of session

17:00 Pub Quiz

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Break out rooms - meet the speakers



Session VII - 28 august 2020 Moderator: Lucie Delemotte

09:00 Deciphering modulation of ligand-gated ion channels Erik Lindahl, Stockholm University 09:20 Understanding the biophysical principles of immune signalling Erdinc Sezgin, Karolinska Institutet Life cell imaging at the nanoscale 09:40 Ilaria Testa, KTH 10:00 Mechanisms that underlie the initiation and resolution of inflammatory bowel disease Eduardo Villablanca, Karolinska Institutet

10:20

10:35

Session VIII Moderator: Emma Lundberg

Coffee Break

10:50 Navigating quantitative proteomics-based networks to gain knowledge on proteome biology in cancer Janne Lehtiö, Karolinska Institutet 11:10 Transcriptional landscapes in health and disease using AI for tissue images and transcriptomes Joakim Lundeberg, KTH 11:30 High resolution anatomical mapping of gene expression using targeted in situ sequencing Mats Nilsson, Stockholm University 11:50 Error correction and alignment of long-read transcriptome sequencing data Kristoffer Sahlin, Stockholm University 12:10 Break out rooms - meet the speakers 12:25 Short break

Session IX

Moderators: Marcel Tarbier & Vaishnovi Sekar		
12:30	RIF-Seq (RNA isolation free sequencing): an innovative platform integrating RNA-seq with high throughput compound screening Tomasz Krzywkowski, Stockholm University	
12:40	The role of entropy in the evolutionary history of proteins captured by Ancestral Sequence Reconstruction David Hueting, KTH	
12:50	The impact of mobility patterns on COVID-19 infection rates Patrick Bryant, Stockholm University	
13:00	Break out rooms - meet the speakers	
13:15	Closing note from SciLifeLab Visualizing research at SciLifeLab Campus Solna - Digital Vernissage & Winner announcement	

Olli Kallioniemi, Director SciLifeLab