

Monday 21 Oct.	
9:00 - 10:00	Registration and welcome + coffee
10:00 - 10:05	Opening address
10:05 - 11:00	Christina Leslie Decoding epigenomic programs in cancer and immunity
11:00 - 11:30	Nikolay Oskolkov National Bioinformatics Infrastructure Sweden (NBIS)
11:30 - 12:00	Christoph Börlin Modeling TF binding profiles through competitive binding
12:00 - 12:30	Fredrik Karlsson Bioinformatics and machine learning for generation and characterization of advanced cell models for drug discovery
12:30 - 13:30	Lunch
13:30 - 14:30	Lior Pachter TBA
14:30 - 15:00	Aleksej Zelezniak TBA
15:00 - 16:00	Coffee break + Poster session 1
16:00 - 18:00	Anna Reymer Workshop TBA
	Johan Bengtsson-Palme Workshop Wait a minute! – On annotation errors and how to spot them
18:00 - 19:30	Dinner (at Wallenberg Conference Center)

Tuesday 22 Oct.	
9:00 - 10:00	Mihaela Zavolan Learning miRNA-dependent regulatory networks from high-throughput data
10:00 - 10:30	Sara Younes Identification of combinatorial markers in Systemic Lupus Erythematosus and disease subtypes
10:30 - 11:00	Coffee break + Poster session 2
11:00 - 11:30	Olukayode Daramola Differential Expression of Annotated Adenylate Cyclase Genes in various Life Stages of <i>Fasciola hepatica</i>
11:30 - 12:00	Marcela Davila University of Gothenburg Bioinformatics Core Facility
12:00 - 12:30	Johanna Hörberg Regulatory role of torsional stress on bZIP transcription factors Interactions with DNA
12:30 - 13:30	Lunch
13:30 - 14:00	Francesco Gatto Elypta
14:00 - 14:30	Erik Larsson Lekholm TBA
14:30 - 15:00	Coffee break
15:00 - 16:00	Careers in Computational Biology (hosted by RSG-Sweden) [http://www.rsg-sweden.iscbsc.org/]
16:00 - 16:05	Closing remarks

Legend

- Keynote speaker
- Sponsor presentation
- Invited speaker
- Oral presentation
- Workshop/Career session
- Poster session