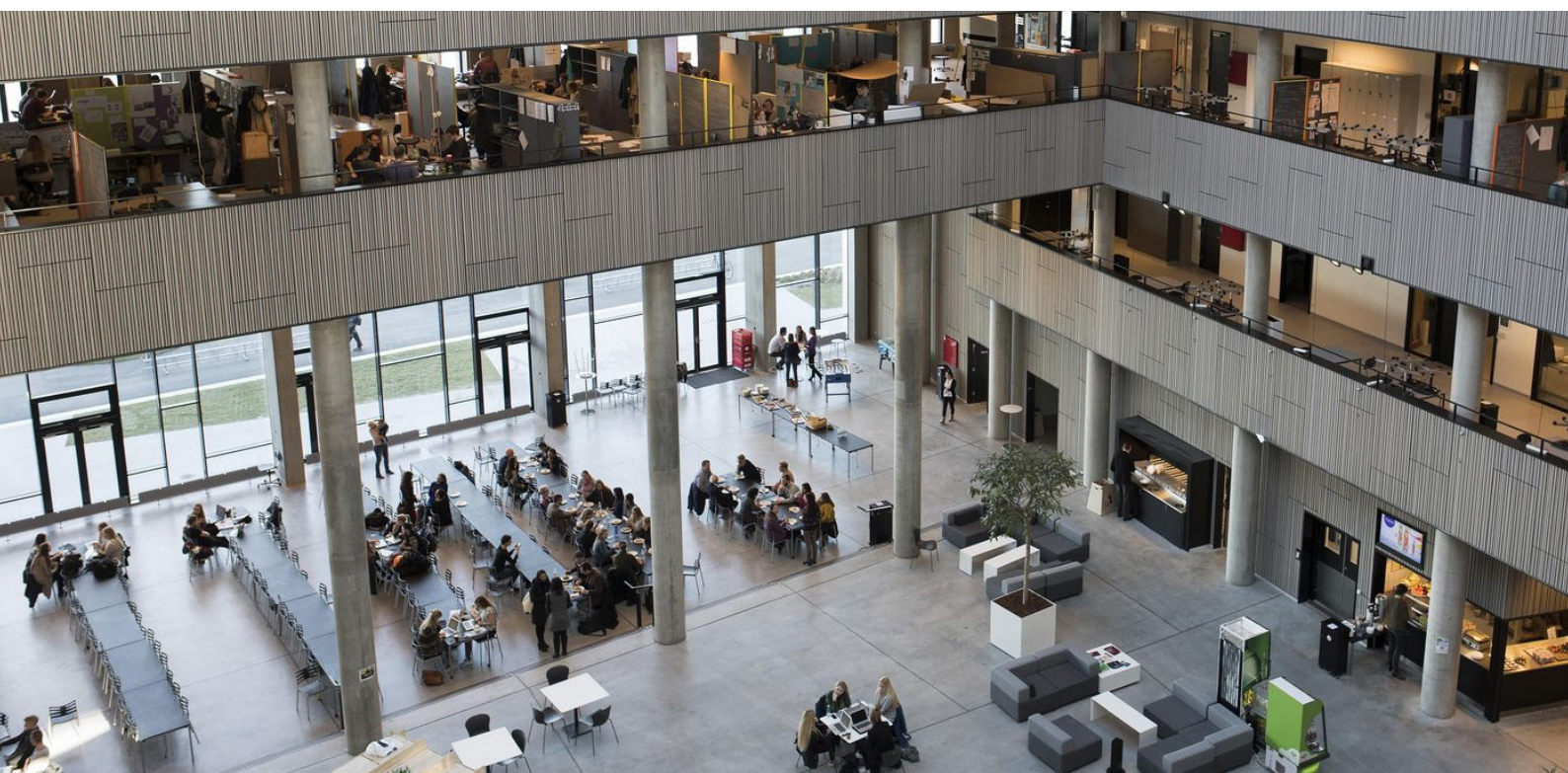




ANNUAL DANISH BIOINFORMATICS CONFERENCE

NOV 18TH &
19TH 2021

CONFERENCE
HANDBOOK



01	KEYNOTE SPEAKERS
06	WORKSHOP OVERVIEW
08	PRACTICAL INFORMATION
10	POSTER SESSION
19	BUSSES TO AALBORG
20	APPENDIX 1 PROGRAMME
23	APPENDIX 2 PARTICIPANT LIST

Sohini Ramachandran

Professor
Brown University

[Google Scholar Page](#)



SESSION

EVOLUTION & POPULATION GENOMICS

ABSTRACT PRESENTATIONS

Reconstructing the evolutionary history of CCR5-Δ32

Leonardo Cobuccio, PhD Student, CPR UCPH

Global faba bean diversity and adaptation

Cathrine Kiel Skovbjerg, PhD Student, MBG AU

SESSION CHAIRS

Thomas Bataillon, Associate Professor, BiRC AU

Palle Villesen, Associate Professor, BiRC AU

Kristian Hanghøj, Postdoc, BIO UCPH

David
Fenyo

Professor
New York University

[Google Scholar Page](#)



SESSION

COMPUTATIONAL MASS SPECTROMETRY & PROTEOMICS

SHORT PRESENTATIONS

MS2AI bridges the gap between MS data and AI applications
Tobias Rehfeldt, PhD Student, IMADA SDU

Quantitative proteomics reveals the causes of bradycardia in heart failure
Konstantin Kahnert, PhD Fellow, BMI UCPH

How to waste money in metabolomics
Johan Lassen, PhD Student, BIRC AU

SESSION CHAIRS

Veit Schwämmle, Associate Professor, BMB SDU
Thomas Koed Doktor, PhD, BMB SDU

Michael Snyder

Professor
Stanford University

[Google Scholar Page](#)



SESSION MACHINE AND DEEP LEARNING

SHORT PRESENTATIONS

Variational Autoencoders to decipher genetic ancestry
Felix Pacheco, MSc Student, CPR UCPH

The national health data science sandbox for training and research
Jennifer Anne Bartell, PhD, Sandbox Coordinator, HeaDS UCPH

SESSION CHAIRS

Tugce Karaderi, Assistant Professor, HeaDS UCPH

Matthias Mann

Professor
University of Copenhagen/
Max-Planck Institute

[Google Scholar Page](#)



SESSION

PERSONALISED MEDICINE & SYSTEMS BIOLOGY

SHORT PRESENTATIONS

Efficient, high-resolution bioinformatics approaches for integrative sequencing analyses for personalized medicine

Shilpa Garg, Assistant Professor, BIO UCPH

The PERSIMUNE Gut Profiler (PGP) – a new tool to profile the microbiome including gut metabolic module (GMM) profiles

Mette Jørgensen, Scientist, Rigshospitalet

Exploring immunopathological mechanisms in rheumatic diseases using Omics data integration

Christopher Aboo, PhD Fellow, HST AAU

SESSION CHAIRS

Allan Stensballe, Associate Professor, HST AAU

Christopher Aboo, PhD Fellow, HST AAU

Mark Robinson

Professor
University of Zurich

[Google Scholar Page](#)



SESSION

RNA BIOLOGY & TRANSCRIPTOMICS

SHORT PRESENTATIONS

Unveiling the mechanisms of m6A-YTH RNA-binding proteins in Arabidopsis thaliana and their mRNA target identity

Carlotta Porcelli, PhD Fellow, BIO UCPH

Single-Cell Dynamics of Interacting Mesenchymal and Leukocyte Populations in Advanced NASH

Sofie Bendixen, PhD Fellow, BMB SDU

SESSION CHAIRS

Elena Papaleo, Associate Professor, CPR UCPH

Stefan Seemann, Associate Professor, IVH UCPH

WORKSHOP SESSION 1

MULTIOMICS DATA INTERGRATION WITH MIXOMICS

Christopher Aboo/Sebastien Dejean

This workshop will explore how a more holistic view of a system can be attained, by combining multiple 'omics analyses. This talk will present the mixOmics R package that proposes multivariate statistical methods to explore and integrate 'omics data sets.

FAIRNESS AND BIAS IN MACHINE LEARNING

Tibor Varga

This workshop will introduce the fundamentals of algorithmic bias and fairness. Learn a few tips and tricks on deep learning tools for biological data: their strengths and shortcomings, how to properly develop them, and how to deploy them to maximise their scientific impact.

INTRODUCTION TO DEEP LEARNING TOOLS

Paolo Marcatili

In this workshop we will discuss dataset creation and curation, overfitting and generalisation, data leakage, new architectures, deployment, and maintenance, through a few specific examples.

INTRODUCTION TO UCLOUD

Emiliano Molinaro

The UCloud platform is a Danish resource for cloud computing. This hands-on workshop will present and introduce users to SDU's interactive research environment. For those already familiar with UCloud, we don't recommend this workshop.

WORKSHOP SESSION 2

ELIXIR BIO.TOOLS

Hans Ienasescu/Veit Schwämmle

Bio.tools is the ELIXIR registry of over 22.000 software tools in the life sciences. The participants will actively learn about the data concepts in bio.tools and how the tool platform interlinks with other data initiatives.

CRISPR BIOINFORMATICS

Jan Gorodkin

In the workshop we will introduce computational methods and their principles for CRISPR/Cas9 on-and off-target design and a computational package for on-and off-target post editing verification for which RNA-seq is used to inspect transcriptome variations.

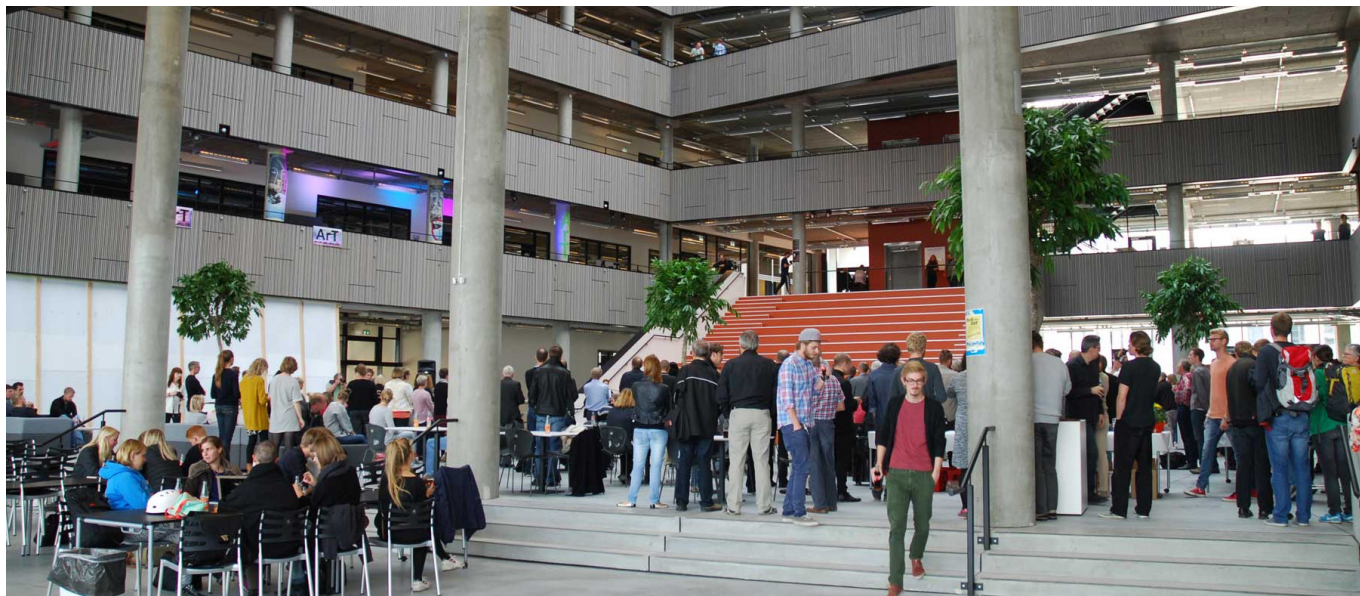
INTRODUCTION TO COMPUTEROME

Darío Carballido

In this workshop, we introduce you to Computerome's custom services available for bioinformaticians. There will be an opportunity to have an open discussion about the best technical solution and services for your future research projects.

PRACTICAL INFORMATION

CONFERENCE VENUE



CREATE CAMPUS AALBORG UNIVERSITY
[RENSBURGGADE 14, 9000 AALBORG](#)

DINNER VENUE



RESTAURANT FUSION
[STRANDVEJEN 4, 9000 AALBORG](#)

POSTER SESSION

POSTER SESSION

For this year's poster session, the preferred format is A1 portrait. Between 12.00-12.50 on Day 1, you will be able to hang your poster in the CREATE Campus Foyer. Poster boards and pins will be provided.

You will then be able to vote for your favourite poster over the course of the conference. Voting will close at 13.00 on Day 2.

Votes will then be counted and the winner announced during the closing ceremony at 15.30. The prize for the best poster is 2000 DKK.

For where to hang your poster, on the next page you can find your name and corresponding number.



POSTER SESSION

- 1 **Adrian Gabriel Zucco**
Personalized survival probabilities for SARS-CoV-2 positive patients by explainable machine learning
- 2 **Adrian Geissler**
Stress: Bacillus subtilis' secretion-life balance in its co-expression network
- 3 **Alba Refoyo Martínez**
Reconstructing ancient HLA haplotypes, using variation graphs and pangenome assemblies
- 4 **Alexander G. B. Grønning**
MultiPep: A hierarchical deep learning approach for multi-label classification of peptide bioactivities
- 5 **Alexander Gmeiner**
Whole Genome Sequencing and Machine Learning to Improve Food Safety
- 6 **Alexander Hauser**
Personalised medicine through GPCR pharmacogenomics
- 7 **Amalia Bogri**
The effect of antimicrobial treatments and environmental stresses on the gut microbiome of cockroaches: a compositional metagenomic analysis
- 8 **Andreas Abildskov Thomsen**
Cancer Research by MALDI Mass Spectrometry Imaging Technology
- 9 **Andreas Møller**
Identifying regulatory networks governing cellular identity in health and disease

POSTER SESSION

- 10 **Anne Hartebrødt**
Principal Component Analysis for Federated Bioinformatic Applications
- 11 **April Snøfrid Kleppe**
Contributing mutagenic processes of the human genomic diversity
- 12 **Arthur Grimaud**
Quantification of proteoforms from the MS/MS fragment ions of large peptides in middle-down proteomics
- 13 **Asta M. Rasmussen**
Characterization of Circular Stable Intronic RNAs in Bladder Cancer
- 14 **Astrid Brix Saksager**
An automatized workflow to study mechanistic indicators for driver gene prediction with Moonlight
- 15 **Astrid Filt**
Detection of sample-swaps in a personalized neoepitope-directed immunotherapy bioinformatics pipeline
- 16 **Carlotta Porcelli**
Unveiling the mechanisms of m6A-YTH RNA-binding proteins in Arabidopsis thaliana and their mRNA target identity
- 17 **Cas Blaauw**
O-GalNAc protein glycosylation prediction using protein language models

POSTER SESSION

- 18 **Catarina Seidi**
A Comprehensive Bioinformatics Analysis of Male Germ Cell Gene Signatures as Putative Oncogenic Biomarkers
- 19 **Derya Aytan-Aktug**
Prediction of plasmid host ranges using random forest
- 20 **Emil Aarre Sørensen**
Microflora Danica: Single primer enrichment of bacterial rRNA genes
- 21 **Erik Fogh Sørensen**
Sex-specific population structure in baboons revealed by hundreds of complete X-chromosomes
- 22 **Eva Harvald**
Delineating transcriptional network in proliferating cardiomyocytes during heart development
- 23 **Felix Teufel**
SignalP 6.0 achieves signal peptide prediction across all types using protein language models
- 24 **Frederik Teudt**
Visualization of Plasmid Diversity in Sewage
- 25 **Gisle Vestergaard**
MAGinator: Combining the best from contig and gene-based metagenomics binning for sensitive and accurate abundance profiling and strain-resolution genomics
- 26 **Gustav Alexander Poulsgaard**
Sequence dependencies and rate estimates of localized mutational processes in cancer

POSTER SESSION

- 27 **Hannah-Marie Martiny**
Identifying genetic signatures in regions around antimicrobial resistance genes
- 28 **Henrietta Holze**
Extending the STRING database with host-parasite protein-protein interactions
- 29 **Iker Rivas-González**
Pervasive incomplete lineage sorting in primates and its genomic and functional determinants
- 30 **Jilong Ma**
Evolution of the Sociality using Social Spiders from Genus Stegodyphus as a Model Organism
- 31 **Johanne Ahrenfeldt**
Temporal dissection of altered pathways during the evolution of cancer
- 32 **Johannes Hadsund**
Positional commonalities of B-cell receptor across independent sequencing studies
- 33 **Judit Szarvas**
Phylogenetic dashboard in the European Bioinformatics Institute's COVID-19 Data Portal
- 34 **Katerina Nastou**
Can dictionary-based text mining be improved with deep learning?

POSTER SESSION

- 35 **Katia Bougiouri**
Inferring patterns of natural selection in ancient Siberian dog lineages
- 36 **Kirstine Rasmussen**
Pathway-resolved investigation of metabolomics data
- 37 **Knud Nor Nielsen**
Structural variant detection using pangenome graphs - a case study with an entomopathogenic fungus
- 38 **Kristian Andersen**
Identifying the transcriptional profile of cardiomyocytes in the regenerating zebrafish heart and translating this to mouse and human
- 39 **Kristoffer Vitting-Seerup**
Isoforms are an overlooked challenge and opportunity in chimeric antigen receptor cell therapy
- 40 **Laura Sans Comerma**
Generating protein representations from tertiary structure. An application of deep learning for protein-protein prediction
- 41 **Lavinia Ioana Fechete**
Transcriptomic response of white clover to cold stress
- 42 **Liubov Pashkova**
The effect of CpG methylation on the CRISPR-Cas9 off-target activity

POSTER SESSION

- 43 **Ludvig Renbo Olsen**
Unsupervised learning of transcription factor binding patterns enables accurate detection and discrimination of different cancer types from shallow whole-genome sequencing of cell-free DNA
- 44 **Marianne Helenius**
White blood cell count variations in acute lymphoblastic leukemia: a COST LEGEND genotype-phenotype study
- 45 **Marie Louise Jespersen**
Global phylogeny of bacterial strains from sewage suggests that local selection shapes geographical phylogenetic clustering
- 46 **Mario Garcia Urena**
Uncovering genes that interact with physical activity in their influence on cardiovascular disease risk
- 47 **Markus Hodal Drag**
Nanopore sequencing of circulating cell-free DNA from blood reveals differentially methylated regions in lean and obese Göttingen Minipigs
- 48 **Mathias Bøgebjerg**
Active transitivity clustering
- 49 **Mathias Stokholm**
GEDI: an R package to enable transcriptomic data integration
- 50 **Matteo Tiberti**
Collection and annotation of cancer-related protein variants with CancerMuts

POSTER SESSION

- 51 **Matthias Mattanovich**
BFAIR: Automated Metabolic Flux Analysis
- 52 **Mette Jørgensen**
The PERSIMUNE Gut Profiler (PGP) – a new tool to profile the microbiome including gut metabolic module (GMM) profiles
- 53 **Mike Krogh Terkelsen**
Spatial mapping of hepatocellular heterogeneity in human NASH
- 54 **Moisès Coll Macià**
spaceNNtime: a neural network to predict the date and sampling location of ancient genomes
- 55 **Nikola Tom**
Cancer bioMarker Prediction Pipeline 2 – CAMPP2, a handy bioconductor package for quantitative data holders
- 56 **Oana Palasca**
Comparison of temporal blood transcriptomes of mouse and pig colitis models and ulcerative colitis patients
- 57 **Patrick Munk**
Global sewage metagenomics provides unparalleled insight into spatial, taxonomic, and genomic evolution of antimicrobial resistance
- 58 **Paula Tataru**
Automatic cell type prediction for scRNA-seq data using QIAGEN CLC Single Cell Analysis Module

POSTER SESSION

- 59 **Peter Husen**
Developing a computational proteomics data sandbox on UCloud
- 60 **Peter Roland Sørensen**
Single-cell dynamics of mesenchymal-leukocyte interactions in advanced NASH
- 61 **Preston Leung**
Polygenic Interactive Discovery System – A tool for exploring the heritability of complex disease as a function of polygenic risk and biological pathways
- 62 **Ragnhild Laursen**
Robust estimation of mutational signatures using a flexible parametrization
- 63 **Ramtin Zargari Marandi**
A machine learning pipeline based on the gut microbiome data for pre-transplant prediction of acute graft-vs-host disease in allogeneic hematopoietic stem cell transplant recipients
- 64 **Rasa Muktupavela**
Modelling the spatiotemporal spread of beneficial alleles using ancient genomes
- 65 **Rebecca Kirsch**
Physical Interaction Networks in STRING v11.5
- 66 **Steffimol Rose**
n/a

POSTER SESSION

- 67 **Thomas Bygh Nymann Jensen**
MICROFLORA DANICA - The Microbiome of Denmark
- 68 **Tobias Frisch**
FeatureCloud - Federated learning on biomedical data
- 69 **Tobias Hegelund Olsen**
Deep learnt antibody representations using vast B-cell repertoires
- 70 **Trine Zachariassen**
Identifying optimal signature genes de novo enabling strain-level resolution and profiling of the microbiome
- 71 **Tue Bjerg Bennike**
The newborn innate immune system development during the first week of life deciphered by high-throughput plasma proteomics
- 72 **Xiaoyu Zhou**
RNA isoform variation detected by combining the advantages of scRNA and nanopore sequencing
- 73 **Xueer Han**
Towards identifying functional SNPs in structured regions of long non-coding RNAs
- 74 **Ying Sun**
Heterogeneity of Topologically Associating Domain (TAD) boundaries by Sequence Information

GETTING TO AALBORG BY BUS

FROM COPENHAGEN

Contact Person
Hiroki Takano/81 90 64 45

LOCATION

[Ingerslevsgade, 1704 København](#)

THURSDAY 18TH OUTBOUND

Departure: 04.30
Arrival: 10.45

FRIDAY 19TH RETURN

Departure: 17.00
Arrival: 23.00

FROM ODENSE

Contact Person
Myhanh Nguyen/51 64 74 92

LOCATION

[Odense Rutebilstation, Dannebrogsgade 10, 5000 Odense](#)

THURSDAY 18TH OUTBOUND

Departure: 06.45
Arrival: 10.45

FRIDAY 19TH RETURN

Departure: 17.00
Arrival: 21.00

FROM AARHUS

Contact Person
Myhanh Nguyen/51 64 74 92

LOCATION

[Thomas Jensens Allé 2, 8000 Aarhus](#)

THURSDAY 18TH OUTBOUND

Departure: 08.45
Arrival: 10.45

FRIDAY 19TH RETURN

Departure: 17.00
Arrival: 19.00

ANNUAL DANISH BIOINFORMATICS CONFERENCE

NOV 18TH &
19TH 2021

CONFERENCE
PROGRAMME

THURSDAY 18TH NOVEMBER DAY 1

TIME/AREA

ACTIVITY

10.30 - 11.00

CREATE CAMPUS FOYER

PARTICIPANT CHECK-IN & MORNING COFFEE

11.00 - 12.00

WORKSHOP SESSION 1

AUDITORIUM 3.107

MULTIOMICS DATA INTERGRATION WITH MIXOMICS
Christopher Abou/Sebastien Dejean

RBG14 5.127

FAIRNESS & BIAS IN MACHINE LEARNING
Tibor Varga

RBG14 5.125

INTRODUCTION TO DEEP LEARNING TOOLS
Paolo Marcatili

RBG14.4/4.513

INTRODUCTION TO UCLLOUD
Emiliano Molinaro

12:00 - 12:50

CREATE CAMPUS FOYER

LUNCH & POSTER HANGING

12:50 - 13:00

AUDITORIUM 3.107

CONFERENCE OPENING
Allan Stensballe & Veit Schwämmle

13:00 - 14:15

AUDITORIUM 3.107

SESSION: EVOLUTION & POPULATION GENOMICS
Keynote: Sohini Ramachandran

ANNUAL DANISH BIOINFORMATICS CONFERENCE

NOV 18TH &
19TH 2021

CONFERENCE
PROGRAMME

THURSDAY 18TH NOVEMBER DAY 1

TIME/AREA

ACTIVITY

14:15 - 14:45

CREATE CAMPUS FOYER

COFFEE BREAK & POSTER SESSION

14:45 - 16:00

AUDITORIUM 3.107

SESSION: COMPUTATIONAL MASS SPECTROMETRY
& PROTEOMICS

Keynote: David Fenyo

16:00 - 17:15

AUDITORIUM 3.107

SESSION: MACHINE & DEEP LEARNING

Keynote: Michael Snyder

17:15 - 17:30

CREATE CAMPUS FOYER

COFFEE BREAK

17.30 - 18.30

RBG14 5.125

WORKSHOP SESSION 2

ELIXIR BIO.TOOLS

Hans Ienasescu/Veit Schwämmle

AUDITORIUM 3.107

CRISPR BIOINFORMATICS

Jan Gorodkin

RBG14 5.127

INTRODUCTION TO COMPUTEROME

Darío Carballido

19:30

STRANDVEJEN 4,
9000 AALBORG

DINNER

RESTAURANT FUSION

ANNUAL DANISH BIOINFORMATICS CONFERENCE

NOV 18TH &
19TH 2021

CONFERENCE
PROGRAMME

FRIDAY 19TH NOVEMBER DAY 2

TIME/AREA

ACTIVITY

09:00 - 10:15
AUDITORIUM 3.107

SESSION: PERSONALISED MEDICINE & SYSTEMS BIOLOGY
Keynote: Matthias Mann

10:15 - 10:45
CREATE CAMPUS FOYER

COFFEE BREAK

10:45 - 11:30
AUDITORIUM 3.107

FUTURE PERSPECTIVES ON COMPUTEROME
Christine Overgaard Rasmussen & Hossein Aghili

11:30 - 13:00
CREATE CAMPUS FOYER

LUNCH & POSTER SESSION

13:00 - 13:30
AUDITORIUM 3.107

BIOINFORMATICS AT THE NATIONAL GENOME CENTER
Ole Lund

13:30 - 14:45
AUDITORIUM 3.107

SESSION: RNA BIOLOGY & TRANSCRIPTOMICS
Keynote: Mark Robinson

14:45 - 15:15
CREATE CAMPUS FOYER

COFFEE BREAK & POSTER REMOVAL

15:15 - 15:30
AUDITORIUM 3.107

CBIOVIKINGS STUDENT ORGANISATION PRESENTATION

15:30 - 16:00
AUDITORIUM 3.107

CONFERENCE CLOSING & POSTER AWARD CEREMONY
Allan Stensballe & Veit Schwämmle

First name	Last name	Institution
Adrian	Otamendi Laspiur	DTU - Technical University of Denmark
Adrian	Geissler	UCPH - University of Copenhagen
Adrian Gabriel	Zucco	UCPH - University of Copenhagen
Alba	Refoyo Martínez	UCPH - University of Copenhagen
Alex Vincent	Thorn	DTU - Technical University of Denmark
Alexander	Gmeiner	DTU - Technical University of Denmark
Alexander	Hauser	UCPH - University of Copenhagen
Alexander G. B.	Grønning	UCPH - University of Copenhagen
Alexandra	Balicki	DTU - Technical University of Denmark
Allan	Stensballe	AAU - Aalborg University
Álvaro	Bustos Gutiérrez	UCPH - University of Copenhagen
Amalia	Bogri	DTU - Technical University of Denmark
Amelie	Fritz	DTU - Technical University of Denmark
Andreas	Møller	SDU - University of Southern Denmark
Andreas Abildskov	Thomsen	SDU - University of Southern Denmark
Anastasiya	Shen	UCPH - University of Copenhagen
Anne	Hartebrodt	SDU - University of Southern Denmark
Annette	Lien	DTU - Technical University of Denmark
April Snøfrid	Kleppe	AU - Aarhus University
Arman	Simonyan	UCPH - University of Copenhagen
Arthur	Grimaud	SDU - University of Southern Denmark
Asbjørn	Kjær	AU - Aarhus University
Ashot	Margaryan	NGC
Asta	M. Rasmussen	AU - Aarhus University
Astrid	Brix Saksager	DTU - Technical University of Denmark
Astrid	Filt	DTU - Technical University of Denmark
Azra	Leto	AAU - Aalborg University
Baptiste	Avot	DTU - Technical University of Denmark
Benjamin	Nichum	AU - Aarhus University
Bent	Petersen	UCPH - University of Copenhagen
Brede	Sørland	UCPH - University of Copenhagen
Camilla	Lemvigh	DTU - Technical University of Denmark
Carina	Jønck	UCPH - University of Copenhagen
Carlota	Carbajo Moral	DTU - Technical University of Denmark
Carlotta	Porcelli	UCPH - University of Copenhagen
Carolina	Barra Quaglia	DTU - Technical University of Denmark
Caroline	Lennartsson	UCPH - University of Copenhagen
Cas	Blaauw	UCPH - University of Copenhagen
Catarina	Seidi	Bangor University
Cathrine	Jespersgaard	NGC
Cathrine Kiel	Skovbjerg	AU - Aarhus University
Catrine	Høm	NGC
Celia	Burgos Sequeros	DTU - Technical University of Denmark
Charles	Vesteghem	Region Nordjylland
Christian	Breinholt	UCPH - University of Copenhagen
Christian	Kronborg	NGC
Christian	Thygesen	Evaxion Biotech
Christina	Wahl Larsen	Aalborg Universitetshospital
Christine	Rasmussen	DTU - Technical University of Denmark
Christopher	Aboo	AAU - Aalborg University

Christopher	Kaae	DTU - Technical University of Denmark
Clara	Drachmann	DTU - Technical University of Denmark
Dario	Carballido	DTU - Technical University of Denmark
Dave Edison	Rojas Calderon	DTU - Technical University of Denmark
Dayana	De Menezes	DTU - Technical University of Denmark
Derya	Aytan-Aktug	DTU - Technical University of Denmark
Dimitrios	Pediotidis Maniatis	AU - Aarhus University
Ditte Caroline	Andersen	SDU - University of Southern Denmark
Ditte Helene	Lundsted	Aalborg Universitetshospital
Doug	Speed	AU - Aarhus University
Dovydas	Vaitkus	UCPH - University of Copenhagen
Elena	Papaleo	DTU - Technical University of Denmark
Ella	Hedeboe	DTU - Technical University of Denmark
Emil Aarre	Sørensen	AAU - Aalborg University
Emiliano	Molinaro	SDU - University of Southern Denmark
Erik	Fogh Sørensen	AU - Aarhus University
Eva	Harvald	SDU - University of Southern Denmark
Felix	Teufel	UCPH - University of Copenhagen
Felix	Pacheco	DTU - Technical University of Denmark
Frederik	Teudt	DTU - Technical University of Denmark
Frederik Nørby Friis	Sørensen	UCPH - University of Copenhagen
G.	Dudnik	Region H
Gabriel	Renaud	DTU - Technical University of Denmark
Georgios-Nikiforos	Pyrounakis	DTU - Technical University of Denmark
Gian-Andrea	Conzett	DTU - Technical University of Denmark
Giorgia	Moranzoni	DTU - Technical University of Denmark
Gisle	Vestergaard	DTU - Technical University of Denmark
Giulia	Corsi	UCPH - University of Copenhagen
Guillaume	Ramstein	AU - Aarhus University
Gül Sude	Demircan	DTU - Technical University of Denmark
Gustav Alexander	Poulsgaard	AU - Aarhus University
Haja	Kadarmideen	Synomics
Hakan	Aktug	RUC - Roskilde University
Hannah-Marie	Martiny	DTU - Technical University of Denmark
Hans	Ienasescu	DTU - Technical University of Denmark
Heidi	Christensen	AAU - Aalborg University
Henrietta	Holze	UCPH - University of Copenhagen
Henrik	Nielsen	DTU - Technical University of Denmark
Henrik H.	De Fine Licht	UCPH - University of Copenhagen
Hiroki	Takano	DTU - Technical University of Denmark
Hossein	Aghili	DTU - Technical University of Denmark
Huijiao	Yang	DTU - Technical University of Denmark
Ib Thorsgaard	Jensen	AU - Aarhus University
Ida	Egendal	Aalborg University Hospital
Iker	Rivas-González	AU - Aarhus University
Ilka	Hoof	LEO Pharma
Iñigo	Prada	UCPH - University of Copenhagen
Ioannis	Louloudis	UCPH - University of Copenhagen
Isabel	Díaz-Pinés Cort	DTU - Technical University of Denmark
Issa Ismail	Issa	AAU - Aalborg University
Jacopo	Vegni	UCPH - University of Copenhagen

Jake	Pham	DTU - Technical University of Denmark
Jakob	Madsen	UCPH - University of Copenhagen
Jamie	Hill	QIAGEN Aarhus
Jan	Gorodkin	UCPH - University of Copenhagen
Jeanette	Bach	AAU - Aalborg University
Jennifer	Bartell	UCPH - University of Copenhagen
Jens Kjærgaard	Boldsen	BioLib
Jeppe	Hallgren	BioLib
Jesper	Christiansen	DTU - Technical University of Denmark
Jilong	Ma	AU - Aarhus University
Johan	Lassen	AU - Aarhus University
Johanne	Ahrenfeldt	AU - Aarhus University
Johannes	Hadsund	SDU - University of Southern Denmark
Jonathan	Robinson	BioInnovation Institute
Jorge	Hernansanz	UCPH - University of Copenhagen
Jørn	Emborg	BioLib
Jose Juan	Almagro Armenteros	UCPH - University of Copenhagen
Josefine	Severin	AAU - Aalborg University
Judit	Szarvas	DTU - Technical University of Denmark
Julian	Regalado	UCPH - University of Copenhagen
Karolina	Krzysińska	DTU - Technical University of Denmark
Kasper	Holm	NGC
Kasper	Munch	AU - Aarhus University
Kat	Steinke	Odense University Hospital
Katerina	Nastou	UCPH - University of Copenhagen
Katia	Bougiouri	UCPH - University of Copenhagen
Katrin	Thomsen	AAU - Aalborg University
Katrine	P. Jensen	AAU - Aalborg University
Kim	Ravnskjær	SDU - University of Southern Denmark
Kirstine	Rasmussen	UCPH - University of Copenhagen
Knud	Nor Nielsen	UCPH - University of Copenhagen
Konstantin	Kahnert	UCPH - University of Copenhagen
Kristian	Andersen	SDU - University of Southern Denmark
Kristina	Koldby	AAU - Aalborg University
Kristine	Degn	DTU - Technical University of Denmark
Kristoffer	Vitting-Seerup	DTU - Technical University of Denmark
Lars	Rønn Olsen	DTU - Technical University of Denmark
Laura	Sans Comerma	DTU - Technical University of Denmark
Laura	Andersen	AU - Aarhus University
Laura	Rose	UCPH - University of Copenhagen
Laura	Skrubbeltrang Hansen	AU - Aarhus University
Laura Johanne	Grønholt	SDU - University of Southern Denmark
Lavinia Ioana	Fechete	AU - Aarhus University
Leonardo	Cobuccio	UCPH - University of Copenhagen
Liubov	Pashkova	UCPH - University of Copenhagen
Lotte	Hindhede	Aarhus University Hospital
Ludvig Renbo	Olsen	AU - Aarhus University
Mads	Brøgger	BioLib
Mads	Pedersen	BioLib
Mads	Sønderkær	Aalborg University hospital
Maja Zimmer	Jakobsen	AAU - Aalborg University

Malene	Revsbech Christiansen	UCPH - University of Copenhagen
Marc	Solé Estragués	AU - Aarhus University
Maria	Speed	AU - Aarhus University
Mariadaria	Ianni-Ravn	UCPH - University of Copenhagen
Mariadaria	Ianni-Ravn	UCPH - University of Copenhagen
Marianne	Helenius	DTU - Technical University of Denmark
Marie Louise	Jespersen	DTU - Technical University of Denmark
Marie Vibeke	Vestergaard	AAU - Aalborg University
Mario	Garcia Urena	UCPH - University of Copenhagen
Mark	Robinson	University of Zurich
Markus Hodal	Drag	UCPH - University of Copenhagen
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