



Programme

LifeTime Opening Conference

May 6-7, 2019

Monday, May 6, 2019

7:30-8:30	Registration and Breakfast
8:30 - 8:45	Welcoming Address Martin Lohse (Scientific Director, MDC) Axel Pries (Dean, Charité – Universitätsmedizin Berlin; – CEO (interim), Berlin Institute of Health BIH) André Le Bivic (Director of the Institute of Biological Sciences, CNRS) Otmar Wiestler (President, Helmholtz Association)
8:45 - 9:15	Introduction by Nikolaus Rajewsky and Geneviève Almouzni (LifeTime coordinators)
9:15 - 11:40	SESSION 1: SINGLE CELL MULTI-OMICS Hosted by WP3 (leads: Giacomo Cavalli and Amos Tanay)
9:15 - 9:35	Amos Tanay , The Weizmann Institute for Sciences, Rehovot, Israel Introduction to WP3: Single cell multi-omics
9:35 - 9:55	Prisca Liberali , FMI – Friedrich-Miescher-Institute for Biomedical Research, Basel, Switzerland Single cells in space and time during development
9:55 - 10:15	Marcelo Nollmann , CNRS – Centre National de la Recherche Scientifique, Montpellier, France Seeing is believing: new breakthroughs in imaging technologies
10:15 - 10:40	Coffee time
10:40 - 11:00	Alexander Van Oudenaarden , Hubrecht Institute, Utrecht, The Netherlands Lineage reconstruction by single-cell sequencing
11:00 - 11:20	Robert Balderas , BD Biosciences, USA Co-creation of innovative tools for Single Cell Multi-omics
11:20 - 11:30	Mats Nilsson , SciLifeLab, Solna, Sweden Spatial cell-maps of molecularly defined cell types using targeted in situ sequencing
11:30 - 11:40	Marlies Vanden Bempt , VIB, Belgium Single cell profiling of different cancer types before and during immune checkpoint therapy
11:40 - 12:10	CROSS TOPICS: INNOVATION, TRAINING Hosted by WP7 and WP10 (leads: Jan Ellenberg , Susan Gasser , Joachim Schultze , Marek Figlerowicz and Dimitris Thanos)

12:10 - 13:30 Lunch time/poster session in parallel.....

13:30 - 15:30 SESSION 2: DATA SCIENCE, AI & MACHINE LEARNING

Hosted by WP4 (leads: **John Marioni, Marc Marti-Renom and Helen Parkinson**)

13:30 - 13:50 John Marioni, EMBL-EBI, Wellcome Genome Campus, Hinxton
Introduction to WP4: Data science, AI & machine learning

13:50 - 14:10 Fabian Theis, Institute of Computational Biology, Helmholtz Center Munich, Germany
Large-scale lineage and latent-space learning in single-cell genomics

14:10 - 14:30 Jean-Philippe Vert, Google, France
Learning representations of single cell data

14:30 - 14:50 Nir Friedman, HUJI – The Hebrew University of Jerusalem, Rehovot, Israel
From factoids to knowledge - how do we connect the dots?

14:50 - 15:10 Anna Kreshuk, EMBL – The European Molecular Biology Laboratory, Heidelberg, Germany
Image analysis with machine learning

15:10 - 15:20 Matthias Becker, DZNE – German Center for Neurodegenerative Diseases, Bonn, Germany
Memory-Driven Computing: A novel architecture for growing single-cell data sets

15:20 - 15:30 Mario Nicodemi, University of Naples, Italy
Machine learning chromatin architecture in single cells by polymer physics

15:30 – 16:00 Coffee time

16:00 - 18:00 SESSION 3: EXPERIMENTAL DISEASE MODELS

Hosted by WP5 (leads: **Jürgen Knoblich, Jean-Christophe Marine and Giuseppe Testa**)

16:00 - 16:20 Giuseppe Testa, IEO – European Institute of Oncology, Milano, Italy
De humani corporis fabrica: the new landscape of experimental models for charting disease dynamics at single cell resolution

16:20 - 16:40 Jürgen Knoblich, IMBA – Institute of Molecular Biotechnology, Vienna, Austria
Modeling human development and disease in Cerebral organoids

16:40 - 17:00 Cédric Blanpain, ULB – Université Libre de Bruxelles, Brussels, Belgium
Deciphering cellular heterogeneity and lineage segregation during embryonic development, tissue homeostasis and tumorigenesis using single cell RNA sequencing

- 17:00 - 17:20** **Paul Vulto**, Mimetas, Netherlands
Organ-on-a-Chip technology as a routine tool in drug discovery and development
- 17:20 - 17:40** **Barbara Treutlein**, ETH Zurich, Switzerland
TBA
- 17:40 - 17:50** **Eleonora Leucci**, Trace PDX platform-LKI, Leuven, Belgium
TRACE & the EurOPDX Research Infrastructure: providing cutting-edge translational cancer research, expertise, & services.
- 17:50 - 18:00** **Sven Nelander**, Uppsala University, Uppsala, Sweden
Therapeutic targets from big data: integrative discovery of treatments for high-risk neuroblastoma
- 18:00 - 18:15** **Short break**.....
- 18:15 - 18:30** **Sabine Oertelt-Prigione**, Radboud University in Nijmegen, Netherlands
The impact of sex differences on biomedical research
- 18:30 - 19:00** **Keynote: Angelika Eggert**, Charité, Berlin, Germany
Creating the basis for tomorrow's cancer precision medicine
- 19:00 - 19:10** Welcoming Address
Steffen Krach, Permanent Secretary for Higher Education and Research, Berlin, Germany
- 19:10** **Reception**

Tuesday, May 7, 2019

- 9:00 - 9:45** **Keynote: Alfonso Valencia**, BSC – Barcelona Supercomputing Center, Barcelona, Spain
From data to models: infrastructures and resources for the challenges of the future medicine
- 9:45 - 10:15** **CROSS TOPICS: ELSI & COMMUNICATION**
Hosted by WP8 and WP9 (leads: Annelien Bredenoord, Maria Elena Torres-Padilla, Leïla Perié, Wolf Reik and Stephan Preibisch)
- 10:15 - 10:45** **Coffee time**.....
- 10:45 - 12:45** **SESSION 4: LIFETIME LAUNCHPAD – DISEASE FOCUS**
Hosted by WP6 (leads: Peter Lichter and Mihai Netea)
- 10:45 - 11:15** **Mihai Netea**, Radboud University Medical Center, Nijmegen, the Netherlands
Jan van Lunzen, ViiV Healthcare, UK
A functional genomics approach to characterize inflammation in disease: the case of HIV infection

- 11:15 - 11:35** **Bart De Strooper**, KU Leuven, VIB, and UK-Dementia research Institute
Single-cell approaches and humanized Alzheimer Disease models:
opportunities for real progress
- 11:35 - 11:55** **Georg Schett**, Friedrich-Alexander University Erlangen-Nürnberg,
Erlangen, Germany
Towards a molecular pathology of chronic inflammatory disease
- 11:55 - 12:15** **Wilko Weichert**, Institute of Pathology, Technical University Munich (TUM),
Munich, Germany
Cancer. On how to attack the emperor of all maladies.
- 12:15 - 12:25** **Céline Vallot**, Institut Curie, Paris, France
Single-cell chromatin profiling reveals heterogeneity of chromatin
landscapes in breast cancer
- 12:25 - 12:35** **Kathrin Kattler**, Saarland University, Saarbrücken, Germany
Spatial transcriptomic and epigenomic maps of human liver: blueprints
for projection of single cell data into hepatic pseudospace
- 12:35 - 12:45** **Claudia Ctordecka**, IMP – Research Institute of Molecular Pathology,
Vienna, Austria
In depth profiling of cardiac spheres to decipher cardiac progenitor
functionality
- 12:45 - 14:15 Lunch time/poster session in parallel.....
- 14:15 - 16:15** **PARALLEL BREAKOUT SESSIONS (WP3, 4, 5 AND 6)**
FROM THE STATE OF THE ART TO LIFETIME OBJECTIVES
- 16:15 - 17:15 Wrap up and closing.....



Conference Information

Date

May 6-7, 2019

Conference Venue

bcc, Alexanderstr. 11, 10178 Berlin

Organisation & Contact

Max-Delbrück-Center for Molecular Medicine in the Helmholtz Association
Robert-Rössle-Straße 10
13125 Berlin

For information about the conference:

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Electronic Programme and Abstract Booklet

Download: lifetime-fetflagship.eu/index.php/program

Poster Presentation Guidelines

Posters will be displayed during the meeting:

on May 6: 12:10-13:30

on May 7: 12:45-14:15

Authors are asked to be present at their poster during the poster sessions. You will find the number of your poster in the electronic programme and abstract book.

Posters should be mounted on May 6, before lunch and removed on May 7 at 17:00 latest.

Internet:

Internet access via Wireless LAN is free of charge in the bcc. The login is "LifeTimeOpening", password: Launch2019.