

Programme

08.15-09.00 Accreditation

09.00-09.20 Ulrich Kalinke

Welcome address "11th TWINCORE Symposium"

09.20-10.00 Yang Li

A big data approach to infection research: translating genetic variation

into immune function

TOPIC: Bacteriology

10.00-10.30 Helle Krogh Johansen

Pseudomonas aeruginosa adaptation in cystic fibrosis airways

10.30-10.40 Susanne Häussler

Parallel evolutionary paths to produce more than one Pseudomonas aeruginosa

biofilm phenotype

10.40-11.10 Coffee Break

TOPIC: RNA viruses

11.10-11.40 **Shirit Einav**

Towards better understanding and predicting severe dengue: from single cell to

multi-cohort transcriptomic analyses

11.40-11.50 Thomas Pietschmann

High resolution profiling of human antibody responses to Hepatitis C virus

11.50-12.00 Gisa Gerold

Quantitative high-resolution proteomics reveals human pathogenic virus host factors

TOPIC: DNA viruses

12.00-12.30 Florian Erhard

Revealing core features of CMV-induced regulation in single cells

by scSLAM-seq

12.30-12.40 Ulrich Kalinke

Myeloid cells in cytomegalovirus dissemination

12.40-13.40 Lunch

13.40-14.10

TOPIC: Epigenetics, immunometabolism

Stacy Horner

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	Regulation of the epitranscriptome during viral infection
14.10-14.40	Chengjian Xu Epigenetic regulation in allergic disease and infection
14.40-14.50	Matthias Lochner Targeting the cellular fatty acid metabolism in intestinal infection and inflammation

14.50-15.20 Coffee Break

TOPIC: Clinical infectiology

15.20-15.50	Markus Cornberg Impact of HCV cure on the immune system
15.50-16.20	Juan Carlos López-Alfonso A machine learning approach to predict virological relapse in chronic hepatitis B virus infection
16.20-16.30	Frank Pessler Cerebrospinal fluid metabolites as biomarkers to distinguish between viral and autoimmune encephalitis
16.30-16.40	Gérard Krause

Digital surveillance, outbreak response management and analysis system

17.00 Barbecue