



Late Summer School 2025

Tumor Heterogeneity, Evolutionary Dynamics and Treatment Response:

Experiments Meet Computation

24 - 26 September 2025

Humboldt Universität Campus Nord, Haus 18, Philippstraße 13 10115 Berlin, Germany (<u>HU Nord</u>)

Wednesday, 24 September 2025		
09:45 – 10:45	LECTURE [Hörsaal 2] "Mathematical oncology and digital twins for clinical oncology" Heiko Enderling, MD Anderson Cancer Center, Houston, USA	
11:00 – 12:00	RESEARCH TALK [Hörsaal 2] "Leveraging single cell technologies to engineer the immune system" Ido Amit, Weizmann Institute of Science, Rehovot, Israel	
12:00 – 12:45	RESEARCH TALK [Hörsaal 2] "Extrachromosomal DNAs (ecDNAs) in cancer" Anton Henssen, Charité – Universitätsmedizin Berlin, Germany	
15:30 – 16:30	RESEARCH TALK [Hörsaal 2] "Integrating patient-specific experimental models, state-of-the-art molecular biology, and large-scale computation to investigate growth principles and treatment options in cancers of the nervous system" Sven Nelander, Uppsala University, Sweden	

Thursday, 25 September 2025

09:00 – 10:00	LECTURE [Hörsaal 2] "The plastic epigenetic network of cancer" Efrat Shema, Weizmann Institute of Science, Rehovot, Israel
10:30 – 11:30	 Short Talks I [Hörsaal 2] Tancredi Massimo Pentimalli: Spatiotemporal dynamics of tumor microenvironment remodeling Adila Apsara: Gut microbiota shape immune responses in radiation-induced lung toxicity Siddharth Annaldasula: Deconvoluting the tumor microenvironment using somatic mutations
11:30 – 12:30	LECTURE [Hörsaal 2] "Single-cell tumor phylogenetics" Niko Beerenwinkel, ETH Zürich, Switzerland





13:30 – 14:30

RESEARCH TALK [Hörsaal 2]

"Single-molecule and single-cell epigenetics: Decoding the epigenome for cancer research and diagnosis"

Efrat Shema, Weizmann Institute of Science, Rehovot, Israel

17:00 – 18:00

LECTURE [MDC-BIMSB*]

"Quantifying Somatic Evolution in Cancer"

Nicholas McGranahan, University College London, UK

Friday, 26 September 2025

,	
09:00 – 10:00	RESEARCH TALK [Hörsaal 2] "Somatic Evolution, Metastasis and the Impact of Cancer Treatment" Nicholas McGranahan, University College London, UK
10:30 – 11:30	 Vuk Dinovic: Radiotherapy-induced tumor and stromal cell state regulation in PDAC Jonas Berger: Long-read single-cell sequencing reveals highly variable aberrant splicing landscape in spliceosome mutated acute myeloid leukemia Kai Wollek: Computational analysis of T cell receptor repertoires
11:30 – 12:30	RESEARCH TALK [Hörsaal 2] "PRISM - Personalizing Radiotherapy with Integrated Scientific Modeling " Heiko Enderling, MD Anderson Cancer Center, Houston, USA
13:30 – 14:30	RESEARCH TALK [Hörsaal 2] "Modeling tumor progression from single-cell sequencing data" Niko Beerenwinkel, ETH Zürich, Switzerland