

The 3rd MOPRED Symposium 2025

Integrative Genomics and Machine Learning for Precision Health

I - Genomic Embeddings, Knowledge Graphs & Personalized Prediction

10:10 - Moritz Sturm

Genetic risk-aware embeddings of whole-genome data for multimodal applications

10:30 - Julia Gehrmann

Designing a reusable pipeline for integrating tabular data with neuroimaging

10:50 - Arber Qoku

Knowledge-guided matrix factorization for multi-omic perturbation analysis

11:10 - Gerd Specht

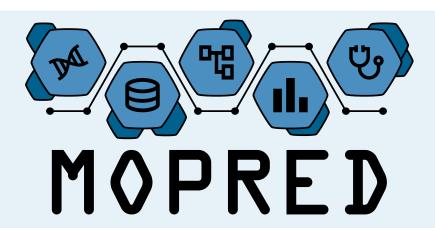
Machine learning approaches for activity-specific health profiling using wearable sensor data

11:30 - Simeon Platte

ML-based molecular burden scores: Predicting treatment outcome based on the genetic variation of underlying mechanisms

Poster Highlights & Networking

- 11:50 Poster Short Talks
- 9 12:00 Group Photo
- 9 12:05 Lunch Break
- 2 12:30 Lunch & Poster Session



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II - Transcriptome Models

13:45 - Jonas Schuck

NanoAIRR: full-length adaptive immune receptor profiling from Nanopore long-read sequencing

14:05 - Nikoletta Katsaouni

Mechanism of action-aware drug ranking with contrastive learning and gene expression profiles

14:25 - Laura Rumpf

Predicting gene-specific regulation with transcriptomic and epigenetic single-cell data

- 14:45 Ewa Szczurek Keynote Lecture Hidden Dimensions of Biology: Latent Models for Multimodal Integration
- 15:25 Poster Session II & Coffee Break

III - Aging, Cancer Biology & Single-Cell Atlases

16:30 – Fabian Kern

Identifying key cellular and molecular aging features in mouse adipose tissue using spatiotemporal transcriptomics

16:50 - Julia Eliseeva

Single-cell RNA sequencing analysis of premalignant lesions to elucidate early cancer progression

17:10 - Christina Kalk

Prediction and validation of split open reading frames across cell types

17:30 - Michael Tyler

Mapping cellular heterogeneity in cancer with the Curated Cancer Cell Atlas

17:55 - Award Ceremony & Closing Remarks

19:00 Dinner (self-funded)