

# Statistical Computing 2018

## 50<sup>th</sup> Workshop - Program Overview

2018-07-08 - 2018-07-11, Schloss Reisenburg (Günzburg)

2018-07-08		2018-07-09	2018-07-10	2018-07-11
	09:20 09:40	<b>Marius Greiff (Dortmund)</b> Optimization of stochastic functions by optimizing expectation and variance	<b>Roman Hornung (München)</b> Random forests for multi-omics data	<b>Sebastian Krey (Köln)</b> Tensile Stress Monitoring of Fibre-Reinforced Plastic
	09:40 10:00	<b>Moritz Berger (Bonn)</b> Tree-Structured Modeling of Time-Varying Coefficients for Discrete Time-to-Event Data	<b>Jörn Lötsch (Frankfurt am Main)</b> A machine-learning based approach to the association of next-generation sequencing derived genotypes with (pain-related) phenotypes	<b>Tobias Hepp (Bonn)</b> Dynamic reference intervals from contaminated data sources
	10:00 10:20	<b>Hryhorii Chereda (Göttingen)</b> Graph-based Convolutional Neural Networks for analyzing pathways in cancer	<b>Lisa Schäfer (Ulm)</b> Repetitive structures pass screening for ordinal relationships	<b>Jakob Richter (Dortmund)</b> Selection of Optimal Subgroup Weights for Survival Analysis
	10:20 10:50	<b>Coffee break</b>	<b>Coffee break</b>	<b>Coffee break</b>
	10:50 11:10	Statistical Computing in Germany : The early years of "Arbeitsgruppen der Arbeitsgruppen" and the flair and impact of Reisenburg	<b>Rolf Backofen (Freiburg)</b>  Computational Genomics: Informatics Problems and Applications in Life Sciences	<b>Thomas Welchowski (Bonn)</b> Correlation-Adjusted Regression Survival Scores for High-Dimensional Variable Selection
	11:10 11:30			<b>Dominic Edelmann (Heidelberg)</b> The distance correlation coefficient for right-censored survival data
	11:30 11:50			
	12:00 13:30	<b>Lunch</b>	<b>Lunch</b>	<b>Lunch</b>
	13:40 14:20	<b>Jennifer Pohle (Bielefeld)</b> State architectures in hidden Markov models	<b>Maike Hohberg (Göttingen)</b> Mixed ordered-continuous copula GAMLSS with an application to poverty dimensions	<b>Departure</b>
	14:20 14:40	<b>Colin Griesbach (Erlangen)</b> New insights in how to boost joint models for longitudinal and time-to-event outcomes	<b>Daniel Schalk (München)</b> compboost: A Modular Framework for Component-Wise Boosting in R	
	14:40 15:00	<b>Anja Rappl (Erlangen)</b> More than one way: Exploring the capabilities of different estimation approaches to joint models.	<b>Gunter Ritter (Passau)</b> Favorite solutions for normal mixture models	
	15:00 15:30	<b>Coffee break</b>	<b>Coffee break</b>	
	15:30 15:50	<b>Andrea Bommert (Dortmund)</b> Benchmark for Filter Methods for Feature Selection in High-dimensional Data	<b>Malte Jastrow (Dortmund)</b> Multimodal likelihood functions occurring in mixture modeling	
	15:50 16:10	<b>Robin Szekely (Ulm)</b> Utilizing foreign classes for feature selection	<b>Marcus Vollmer (Greifswald)</b> Randomized Stepwise Regression – a Modification of Stepwise Methods in Generalized Linear Models and its Application on Sepsis Data	
	16:10 16:30	<b>Giuseppe Casalicchio (München)</b> Visualizing the Feature Importance for Black Box Models	<b>Regina Stegherr (Ulm)</b> Analysing unmeasured baseline covariates in studies with delayed entry using a joint model: fitting the model in presence of left-truncation	
	16:30 16:50	<b>Break</b>	<b>Break</b>	
	16:50 17:50	<b>Phil Bowsher (Indianapolis)</b> An Introduction to Shiny and R Markdown With Applications in Research and Drug Development	<b>Working group meeting</b> on Statistical Computing 2019	
<b>Dinner</b>	18:00 20:00	<b>Dinner</b>	<b>Dinner</b>	
<b>Jan Beyersmann (Ulm)</b> Sampling and Re-Sampling Complex Time-to-Event data: Beyond Standard Cohort and Standard Bootstrap	20:00 21:00	<b>Phil Bowsher (Indianapolis)</b> Hands on Tutorial		