2019 Conference on Lifetime Data Science: Foundations and Frontiers





The 2^{nd} Conference on Lifetime Data Science

University of Pittsburgh Pittsburgh, Pennsylvania, USA May 29 – 31, 2019

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Welcome

On behalf of the Program Committee we are very pleased to extend a warm welcome to the 2019 Conference on Lifetime Data Science: Foundations and Frontiers. This three day event is modeled after the highly successful inaugural lifetime data science conference whose scientific program was co-Chaired by Mei-Cheng Wang and Jack Kalbfleisch and graciously hosted by the University of Connecticut in 2017.

We have another wonderful program this year, beginning with short courses by leading researchers in life history methods which are followed by a welcome reception and poster session taking place from 6:30 to 8:30 p.m. on Wednesday, May 29 at the Wyndham Hotel. We are delighted to have Keynote Presentations by Professors Odd Aalen, Danyu Lin and Ross Prentice, each of whom have made important pioneering contributions in statistical methods for the design and analysis of life history studies. The program is also packed with a series of exciting invited sessions which will make it hard to choose which sessions to attend so we encourage you to plan ahead. This can be achieved using the Whova cell phone app which we encourage you to download if you haven't already!

The many exciting areas of research within lifetime data science are clearly in evidence from our scientific program. The theme this year of "Foundations and Frontiers" was chosen to acknowledge the importance of the foundations of our discipline when tackling new problems arising in the frontiers of modern science and technology. The challenges include high dimensional genomic data, dependent selection and observation schemes, issues surrounding estimands and causal inference, use of administrative data for scientific inquiry, among many others. Statistical science is a young and exciting discipline and we are fortunate that many of the founding researchers continue to be active making important contributions to our discipline and maintain their scientific leadership. You may well find yourself sitting next to such a person at this meeting!

To put a conference together of this magnitude is no small task and there are many people to thank. First we wish to thank Professor Ying Ding and Professor Yu Cheng along with all of the members of the Local Arrangements Committee for their leadership, hard work, and dedication. Their tireless efforts at managing the registration, Whova app, website and facilities have ensured things run smoothly and that we are able to enjoy this wonderful venue. We extend this thanks to the staff devoted to this conference at the University of Pittsburgh and University of Waterloo, and the Department of Biostatistics and Department of Statistics at University of Pittsburgh in particular. We also gratefully acknowledge the Executive Committees of the Lifetime Data Analysis Interest Group, and more recently the ASA Section on Lifetime Data Science, for their wisdom and guidance during the planning stages of the event.

The Student Paper Competition drew a remarkable number of submissions which attests to the health of our discipline. We thank Professor Guoqing Diao and the members of the Student Paper Award Committee for promoting the event, reviewing the many papers, and taking on the difficult task of selecting the winners. There was a very strong pool of candidates and we thank all participants, congratulate the winners and look forward to hearing their presentations during the conference. Their awards will be presented at the conference banquet at the Wyndham Hotel on the evening of May 30 at 6:00 p.m..

This conference also serves as a chance to celebrate the creation of our Section on Lifetime Data Science within the American Statistical Association, which took effect in January of this year. The support expressed for the creation of our section was very strong and the enthusiasm is reflected by the engagement of our members in every aspect of the program.

Finally we would like to thank all session organizers whose efforts have ensured our exceptional scientific program, and the speakers whose work form the foundation of this conference.

In conclusion, we hope you enjoy your visit to Pittsburgh and take the time to renew friendships and extend your networks. We hope there are also lots of opportunities to jointly explore current and future research directions and that you have a productive and fun-filled time at this very special conference.

We are privileged to be co-chairs of this important conference and hope you find it a rewarding event.

Richard Cook Co-Chair, Program Committee University of Waterloo Jianwen Cai Co-Chair, Program Committee University of North Carolina at Chapel Hill

Acknowledgements

The ASA Lifetime Data Science (LiDS) section would like to acknowledge the generous support from the following committees and teams. We also gratefully acknowledge our student volunteers.

Scientific Program Committee

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Program Overview

Wednesday May 29

8:30 - 16:30	Short Courses
	Short Course I: Ørnulf Borgan & Sven Ove Samuelsen Two Phase Studies for Lifetime Data
	Short Course II: Hein Putter Dynamic Prediction in Survival Analysis
	Short Course III: Jing Qin Biased Sampling, Left Truncation and Survival Analysis
18:30 - 20:30	Poster Session and Welcome Reception at Wyndham Hotel

Thursday May 30

8:45 - 9:00	Welcome
9:00 - 10:00	Keynote Presentation: Odd O. Aalen Causal Inference for Survival Data, with Emphasis on Mediation Analysis
10:00 - 10:30	Refreshment Break
10:30 - 12:00	 Invited Sessions Ming-Hui Chen: Recent Development of Bayesian Joint Modeling of Time-to-Event and Longitudinal or Spatial Data Yu Cheng: Novel Semi-parametric Models for Complex Survival Data Guoqing Diao: Advances in the Analysis of Composite Endpoints subject to Component-wise Censoring Noorie Hyun: Survival Analysis Methods for Complex Sample Data Torben Martinussen: Causal Conclusions Based on Cox Regression Analysis Edsel A. Peña: Threshold Modeling, Cancer Risk, and Agreement Assessments Jonathan Siegel: Estimands: The First Update to Regulatory Statistical Guidance in 20 Years - The Pharmaceutical Industry Working Group on Estimands in Oncology Ronghui (Lily) Xu: Prediction and Estimation for Complex Survival Data
13:30 – 15:00	 Invited Sessions 9. Chunyan Cai and Yongseok Park: Clinical Trial Design and Data Analysis with Late-Onset Effects 10. Wei Chen: Advanced Analysis Methods for Survival Outcome with High-Dimensional Data 11. Richard Cook: Founders Session on Current Topics in Lifetime Data Analysis 12. Jon Michael Gran: Causal Inference in Life History Analysis 13. Hélène Jacqmin-Gadda: Joint Models for Time-to-Event and Multiple Longitudinal Data or High Dimensional Data 14. Daniel Nevo: Survival Analysis with Error-Prone or Missing Covariate Measurements 15. Douglas Schaubel: Novel Methods for Complex Censored Data
15:00 - 15:30	Refreshment Break
15:30 - 17:00	 Invited Sessions 17. Jianwen Cai: Advanced Statistical Methods for Time to Event Data in Complex Observational Studies 18. Ying Ding: Recent Advances in Complex Bivariate Time-to-Event Data Modeling and Analysis 19. Ruzong Fan: Complex Data Analysis in the Biomedical and Health Research 20. Chiung-Yu Huang: Recent Advances in Survival and Recurrent Event Data Analysis 21. Lei Liu: Survival Analysis Methods for Alzheimer's Disease 22. Ruth Pfeiffer: Mediation Analysis for High-Dimensional Data 23. Rajeshwari Sundaram: Innovative Applications of Joint Modeling 24. Mei-Ling Ting Lee: Applied Stochastic Models for Time-to-Event Data
18:00 - 20:30	Banquet at Wyndham Hotel

8:00 - 9:00	Keynote Presentation: Ross Prentice Regression Models and Multivariate Life Tables	
9:00 - 10:00	Keynote Presentation: Danyu Lin Semiparametric Regression Analysis of Interval-Censored Data	
10:00 - 10:30	Refreshment Break	
10:30 - 12:00	Invited Sessions	
	 25. Paul Albert: Innovative Approaches for Studying the Effects of Chemical Mixtures on Disease Throughout the Life-span 26. Rebecca A. Betensky: Methods for Dependent Truncation 27. Richard Cook: Methods for Lifetime Data Processes Under Intermittent Observation 28. X. Joan Hu: Cross-Sectional Analysis of Life History Data 29. Aiyi Liu: Novel Approaches for Time-to-Event Data 30. Edsel A. Peña: Multiple Testing, Simultaneous and Joint Modeling, and Calibration 31. Mei-Ling Ting Lee: Genomic Applications with Survival Outcomes 32. Mei-Cheng Wang: New Methods for Risk Prediction and Precision Medicine 33. Ronghui (Lily) Xu: Causal Inference for Survival Data 	
13:30 - 15:00	Invited Sessions	
	 34. Guoqing Diao: Session for Student Paper Award Winners (1) 35. Lu Mao: Causal Inference with Time-to-Event Data 36. Hein Putter: Multistate Models as A Framework for Life History Analysis 37. Thomas Scheike: Models and Applications with Recurrent Events 38. Yu Shen: Challenges and Advances of Research in Health Service Studies 39. Rajeshwari Sundaram: Innovative Methods for Assessing Diagnostic Accuracy and Prediction Accuracy 40. Andrew Titman: Extensions to Joint Longitudinal-Survival Modelling 41. Grace Y. Yi: Emerging Issues and Methods on Censored Data 42. Leilei Zeng: Intermittent Observation of Life History Processes 	
15:00 - 15:30	Refreshment Break	
15:30 - 17:00	Invited Sessions	
	 43. Mouna Akacha: Multi-State Models in Practice 44. Guoqing Diao: Session for Student Paper Award Winners (2) 45. Jong Hyeon Jeong: Recent Developments in Statistical Methods on Semi-Competing Risks Data 46. Bin Nan: Survival Analysis with Missing or Mismeasured Data 47. Hua Shen: Recent Advances in the Analysis of Complex Lifetime Data Involving Recurrent Events 48. Xiao Song: Personalized Treatment Selection with Censored Survival Outcome 49. Jianguo (Tony) Sun: Recent Development in the Analysis of Recurrent Event Data 50. Donglin Zeng: Novel Application of Survival Models in Complex Biomedical Studies 	

Causal Inference for Survival Data, with Emphasis on Mediation Analysis

Odd O. Aalen (University of Oslo, Norway)

Overview

Causal inference is becoming an important theme in survival analysis. We discuss causal mediation analyses for survival data and propose an approach based on the additive hazards model. The emphasis is on a dynamic point of view, that is, understanding how the direct and indirect effects develop over time. To define direct and indirect effects in a longitudinal survival setting we take an interventional approach (Didelez, 2018) where treatment is separated into one aspect affecting the mediator and a different aspect affecting survival. In general, this leads to a version of the non-parametric g-formula (Robins, 1986). In the present talk, we demonstrate that combining the g-formula with the additive hazards model and a linear structural equation model for the mediator process results in simple and interpretable expressions for direct and indirect effects in terms of relative survival as well as cumulative hazards. Our results generalise and formalise the method of dynamic path analysis (Fosen et al, 2006; Strohmaier et al, 2015) and also work by Lange and Hansen (2011). An application will be given.

About the Keynote Speaker

Odd O. Aalen is Professor Emeritus of Biostatistics at the University of Oslo and member of the Oslo Centre for Biostatistics and Epidemiology. His doctoral work, completed at the University of California, Berkeley in 1976,



led to a revolution in the field of survival analysis by reformulating it in terms of counting processes thereby enabling the use of martingales for the development of large sample theory. Odd has made many important contributions to the fields of survival and event history analysis including the development of nonparametric methods of estimation, additive hazards models, advancing understanding of the implications of frailty in sampling and analyses, and causal analysis with life history data through dynamic path analysis. Much of this work is reflected in the highly acclaimed 2008 Springer book with Ørnulf Borgan and Håkon Gjessing entitled *Survival and Event History Analysis: A Process Point of View.* His collaborative work has included infectious disease epidemiology, screening for and treatment of cancer, the effects of gestational alcohol exposure, and studies of employment and disability. He is Fellow of the American Statistical Association, was the 2008 Armitage Lecturer at the University of Cambridge and is an elected member of the Norwegian Academy of Science and Letters.

Regression Models and Multivariate Life Tables

Ross Prentice (Fred Hutchinson Cancer Research Center/University of Washington)

Overview

Regression methods that adapt Cox regression to multivariate failure times, on the same or different failure time axes, will be presented. These methods specify Cox-type semiparametric regression models for marginal single and double failure rates, and use estimating functions and empirical process methods, like those developed by Danyu Lin, L.J. Wei and colleagues for marginal single failure hazard rates, for hazard ratio parameter and for baseline hazard rate estimation. Sandwich type variance process estimators are developed for all model parameters, along with a perturbation resampling procedure for complex constructs of modeled parameters. As a byproduct semiparametric estimators of pairwise survivor functions, given covariates that may be evolving in time, are readily obtained from Peano series representations of these survivor functions in terms of marginal single and double failure rates, and corresponding semiparametric estimators of cross ratio and concordance functions are also readily obtained to characterize pairwise dependencies between failure times given covariates. An application to clinical outcome data from a large low-fat dietary intervention trial among post menopausal women will be presented, and some contrast between these approaches and those based on counting process intensity modeling, as well as on frailty and copula modeling, will be provided. This is joint work with Dr. Shanshan Zhao of NIEHS.

About the Keynote Speaker

Ross Prentice is an ongoing member and former Director of the Public Health Sciences Division of the Fred Hutchinson Cancer Research Center, and is Professor of Biostatistics at the University of Washington.



Ross has made substantial and pioneering contributions in biostatistics and epidemiology with an emphasis on issues arising in the analysis of failure time data. These include the development of innovative research designs, methods for the use of surrogate endpoints, and methods for dealing with covariate measurement error. Ross' work has been motivated through problems arising in his long-standing collaborations in genetic epidemiology, nutritional epidemiology, biomarker studies, and cancer research. He served as Principal Investigator of the Clinical Coordinating Center for the Womens Health Initiative from 1992-2011 and continues to serve as Co-PI. He is a Member (1990) of the Institute of Medicine/National Academy of Sciences, and served on its Food and Nutrition Board. Honors include the Marvin Zelen Leadership Award from Harvard University, the Prentice Professorship for Biostatistical Collaboration at the University of Washington, the Research Excellence in Epidemiology and Prevention Award from the American Association for Cancer Research (AACR), the Team Science Award from the AACR, and the COPSS Presidents and R.A. Fisher Lectureship awards.

Semiparametric Regression Analysis of Interval-Censored Data

Danyu Lin (University of North Carolina at Chapel Hill)

Overview

Interval censoring arises frequently in clinical, epidemiological, financial, and sociological studies, where the event or failure of interest is not observed at an exact time point but is rather known to occur within a time interval induced by periodic monitoring. We formulate the effects of potentially time-dependent covariates on the interval-censored failure time through semiparametric regression models, such as the Cox proportional hazards model. We study nonparametric maximum likelihood estimation with an arbitrary number of monitoring times for each study subject. We develop an EM algorithm that involves very simple calculations and converges stably for any dataset, even in the presence of time-dependent covariates. We show that the estimators for the regression parameters are consistent, asymptotically normal, and asymptotic theory to competing risks and multivariate failure time data. Finally, we demonstrate the desirable performance of the proposed numerical and inferential procedures through simulation studies and applications to real medical studies.

About the Keynote Speaker

Danyu Lin is the Dennis Gillings Distinguished Professor of Biostatistics, a Member of the Lineberger Comprehensive Cancer Center, and a member of the Center for AIDS Research. Danyu is a prolific researcher with most of his articles appearing in leading statistical and genetic journals. He has made many fundamental contributions to statistical methods for



the designs and analysis of biomedical studies, especially in the areas of survival analysis and statistical genetics. Danyu also has extensive applied research experience in cancer, cardiovascular diseases, AIDS, and psychiatry. He has received numerous honors and awards, including Fellow of both the American Statistical Association and the Institute of Mathematical Statistics, Mortimer Spiegelman gold medal from the American Public Health Association for outstanding contributions to biostatistics, and George W. Snedecor Award from the Committee of Presidents of Statistical Societies for the development of statistical theory in biometry. He has served on many editorial and advisory boards, including Associate Editor of Biometrika, Journal of the American Statistical Association and Biometrics, Consultant to the FDA, the Site Visit Team to the NCI Biostatistics Branch, and the External Advisory Committee of Fred Hutchinson Cancer Research Center.

Scientific Program

Wednesday May 29

Wednesday May 29, 8:30 – 16:30			
Wednesday	May 29, 8:30 – 16:30	Short Course I	BENEDUM G29
8:30 - 16:30	Ørnulf Borgan (University of O Sven Ove Samuelsen (Universi	slo, Norway) and ty of Oslo, Norway)	
	Two Phase Studies for Lifetime I	Data	
Wednesday	May 29, 8:30 – 16:30	Short Course II	BENEDUM G30
8:30 - 16:30	Hein Putter (Leiden University	Medical Centre, The Netherlands)	
	Dynamic Prediction in Survival A	Analysis	
Wednesday	May 29, 8:30 – 16:30	Short Course III	BENEDUM G31
8:30 - 16:30	Jing Qin (NIH/NIAID)		
	Biased Sampling, Left Truncation	n and Survival Analysis	
Wednesday	May 29, 10:00-10:30	Refreshment break	BENEDUM Lobby
	14:30-15:00	Refreshment break	BENEDUM Lobby

Wednesday May 29, 18:30 - 20:30

Wednesday May 29, 18:30 - 20:30

18:30 – 20:30 Poster Session and Welcome Reception

Xiaotian Gao (University of Pittsburgh) Inference on Mean Quality-adjusted Lifetime Using Joint Models for Continuous Quality of Life Process and Time to Event

Isabelle Weir (Boston University School of Public Health) Multivariate meta-analysis model for the difference in restricted mean survival times

Pål Christie Ryalen (University of Oslo) The additive hazard estimator is consistent for continuous-time marginal structural models

Tao Sun (University of Pittsburgh) Machine learning with GWAS to predict AMD progression

Cynthia Crowson (Mayo Clinic) Creating and error-checking datasets for survival and multi-state models

Elizabeth Atkinson (Mayo Clinic) Calculating Adjusted Survival Curves

Xinjun Wang (University of Pittsburgh) A comparison and assessment of tree-based methods for subgroup identification with time-to-event data

Jimmy Efird (Cooperative Studies Program Epidemiology Center) Asymptotic Distribution of Constrained Left-Tailed Survival Densities

Wyndham Hotel

18:30 – 20:30 Poster Session and Welcome Reception

Myeonggyun Lee (New York University) Empirical comparison of Sub-cohort sampling designs for Breast cancer risk prediction model on the NYU Woman's Health Study (NYUWHS)

Huining Kang (University of New Mexico) A cost-effective approach to high-throughput gene expression analysis under case-cohort design

Sarah Conner (Boston University) Association between body mass index and risk of atrial fibrillation in the Framingham Heart Study:

g-formula method and difference in restricted mean survival times

Andrew Ying (University of California San Diego) Causal Effects on Birth Defects with Missing by Terathanasia

Niklas Maltzahn (Oslo Center for Biostatistics and Epidemiology) Hybrid Landmark estimation of transition probabilities

Yun-Hee Choi (Department of Epidemiology and Biostatistics, University of Western Ontario) Joint nested frailty models for clustered recurrent and terminal events: An application to colonoscopy screening visits and colorectal cancer risks in Lynch Syndrome families

Busola Sanusi (University of North Carolina at Chapel Hill) Nonparametric Estimation of the Joint Distribution of a Survival Time and Mark Variable in the Presence of Dependent Censoring

Adane F. Wogu (University of North Carolina at Chapel Hill) Proportional Subdistribution Hazards Model for Competing Risks in Case-Cohort Studies

Thursday May 30

Thursday May 30, 8:45 – 9:00

Thursday May 30, 8:45-9:00

8:45 - 9:00 Welcome

Thursday May 30, 9:00 – 10:00

Thursday May 30, 9:00 - 10:00

Keynote Presentation: Odd O. Aalen

Chair: Richard Cook (University of Waterloo, Canada)

9:00 - 10:00Odd O. Aalen (University of Oslo, Norway) Causal Inference for Survival Data, with Emphasis on Mediation Analysis

Thursday May 30, 10:00 – 10:30

Thursday May 30, 10:00–10:30

10:00 – 10:30 Refreshment Break

Thursday May 30, 10:30 - 12:00

Thursday May 30, 10:30 – 12:00

Recent Development of Bayesian Joint Modeling of Time-to-Event and Longitudinal or Spatial Data

Chair: Dandan Liu (University of Vanderbilt)

Organizer: Ming-Hui Chen (University of Connecticut)

10:30 - 10:55 Paul Albert (NIH/NCI) A Joint Model Approach for Longitudinal Data with no Time-Zero and Time-To-Event with Competing Risks 10:55 - 11:20**Guanyu Hu** (University of Connecticut) Bayesian Variable Selection for Cox Regression Model with Spatially Varying Coefficients with Applications 11:20 - 11:45Ming-Hui Chen (University of Connecticut) Assessing Importance of Biomarkers: a Bayesian Joint Modeling Approach of Longitudinal and Survival Data with Semicompeting Risks

Thursday May 30, 10:30 – 12:00

Invited Session 2

BENEDUM G31

Novel Semi-parametric Models for Complex Survival Data

Chair and Organizer: Yu Cheng (University of Pittsburgh)

BENEDUM 157

BENEDUM 157

BENEDUM Hall Lobby

BENEDUM G30

Invited Session 1

Keynote Presentation

10:30 - 10:50	Jing Ning (University of Texas MD Anderson Cancer Center) Semiparametric Model and Inference for Bivariate Survival Data Subject to Biased Sampling
10:50 - 11:10	Sunyoung Shin (University of Texas at Dallas) Ensemble Estimation and Variable Selection with Semiparametric Regression Models
11:10 - 11:30	Xianghua Luo (University of Minnesota) Time-dependent Covariates in Recurrent Event Models
11:30 - 11:50	Thomas Scheike (University of Copenhagen, Denmark) Excess Risk in the Matched Cohort Study

Thursday May 30, 10:30 – 12:00

Invited Session 3

BENEDUM G36

Advances in the Analysis of Composite Endpoints subject to Component-wise Censoring

Chair: Qingxia Chen (Vanderbilt University)

Organizer: Guoqing Diao (George Mason University)

- 10:30 10:50 **Junshan Qiu** (FDA) The Win Ratio : On Interpretation and Handling of Ties
- 10:50 11:10 Audrey Boruvka (Roche, Canada) Computation and Applications in Joint Models for Progression and Survival under Componentwise Censoring
- 11:10 11:30 Ingrid Van Keilegom (KU Leuven, Belgium) A Multivariate Normal Regression Model for Survival Data Subject to Different Types of Dependent Censoring
- 11:30 11:50 **Guoqing Diao** (George Mason University) Semiparametric Regression Analysis for Composite Endpoints Subject to Component-Wise Censoring

Thursday May 30, 10:30 – 12:00

Invited Session 4

Invited Session 5

BENEDUM G27

BENEDUM G29

Survival Analysis Methods for Complex Sample Data

Chair: Jaeun Choi (Albert Einstein College of Medicine)

Organizer: Noorie Hyun (Medical College of Wisconsin)

- 10:30 10:55 **Takumi Saegusa** (University of Maryland, College Park) Survival Analysis for Integrated Data from Multiple Sources
- 10:55 11:20 **Noorie Hyun** (Medical College of Wisconsin) Sample-Weighted Semiparametric Models for Competing Risks Data Subject to Left-/Interval Censoring from Electronic Health Records
- 11:20 11:45 Ai (Andy) Ni (Ohio State University) Concordance Measures in Survival Analysis on Survey Data

Thursday May 30, 10:30 - 12:00

Causal Conclusions Based on Cox Regression Analysis

Chair: Ying Ding (University of Pittsburgh)

Organizer: Torben Martinussen (University of Copenhagen, Denmark)

- 10:30 10:55 **Jonathan Bartlett** (University of Bath, UK) Hazard Ratios - What's Different and What's Not
- 10:55 11:20 Morten Valberg (Oslo University Hospital, Norway) Potential Causal Consequences of Observed Proportional Hazards
- 11:20 11:45 **Torben Martinussen** (University of Copenhagen, Denmark) Subtleties in the Interpretation of Hazard Ratios

Thursday May 30, 10:30 – 12:00 Invited Session 6 **BENEDUM 158** Threshold Modeling, Cancer Risk, and Agreement Assessments Chair: Jonathan Race (Ohio State University) Organizer: Edsel A. Peña (University of South Carolina) 10:30 - 10:55 Michael Pennell (Ohio State University) A Bayesian Semiparametric First Hitting Time Model for Latent Fetal Development 10:55 - 11:20 Jung In Kim (NIH/NIEHS) Accounting for Preinvasive Conditions in the Analysis of Invasive Cancer Risk: With Application to Breast Cancer and the Sister Study **AKM Fazlur Rahman** (University of Alabama at Birmingham) 11:20 - 11:45Nonparametric Regression Method for Broad Sense Agreement

Thursday May 30, 10:30 – 12:00 Invited Session 7 BENEDUM 227

Estimands: The First Update to Regulatory Statistical Guidance in 20 Years - The Pharmaceutical Industry Working Group on Estimands in Oncology

Chair: Ulrich Beyer (Roche)

Organizer: Jonathan Siegel (Bayer)

- 10:30 10:55 **Jonathan Siegel** (Bayer) Survival Trial Design Strategies in an Estimands Framework
- 10:55 11:20 Feng Liu (AstraZeneca) Developing Estimands in Oncology Trials : Understand Scientific Questions of Interest
- 11:20 11:45 **Shoubhik Mondal** (Boehringer Ingelheim) Sensitivity Analyses in Estimands in Oncology

Discussant: Anja Schiel (EMA/CHMP)

Thursday May 30, 10:30 – 12:00

Prediction and Estimation for Complex Survival Data

Chair: Andrew Ying (University of California, San Diego) Organizer: Ronghui (Lily) Xu (University of California, San Diego)

 10:30 - 10:55 Yingwei (Paul) Peng (Queen's University, Canada) Measures of Explained Variation under the Cure Model for Survival Data
 10:55 - 11:20 Wenging He (Western University, Canada)

Parametric and Semiparametric Estimation Methods for Survival Data under a Flexible Class of Models

Invited Session 8

11:20 – 11:45 **Yuan Wu** (Duke University) Predictive Accuracy of Markers or Risk Scores for Interval Censored Survival Data

Thursday May 30, 12:00 – 13:30

Thursday May 30, 12:00 – 13:30

12:00 – 13:30 Lunch Break

BENEDUM G28

Thursday May 30, 13:30 – 15:00

Thursday	May 30, 13:30 – 15:00	Invited Session 9	BENEDUM G37
Clinical Tr	ial Design and Data Analysis wi	th Late-Onset Effects	
Chair:	Chi Hyun Lee (University of Mass	achusetts Amherst)	
Organ Pittsb	izer: Chunyan Cai (University of T urgh)	exas Health Science Center at Houston) and	d Yongseok Park (University of
13:30 - 13:	50 Chunyan Cai (University of ² A Bayesian Design for Phase I	Texas Health Science Center at Houston) II Clinical Trials with Late-onset Responses	Based on Multiple Imputation
13:50 - 14:	3:50 – 14:10 Ruitao Lin (University of Texas MD Anderson Cancer Center) Time-to-Event Bayesian Optimal Phase II Trial Design for Cancer Immunotherapy		
14:10 - 14:	30 Mengling Liu (New York Un Design and Analysis of Clinica	iversity) al Trials in the Presence of Delayed Treatm	ent Effect
14:30 - 14:	50 Yongseok Park (University c Designing Cancer Immunothe	of Pittsburgh) rapy Trials with Random Treatment Time-I	Lag Effect
Thursday	May 30, 13:30 – 15:00	Invited Session 10	BENEDUM G31
Advanced	Analysis Methods for Survival (Outcome with High-Dimensional Data	
Chair:	Ruzong Fan (Georgetown Univers	ity Medical Center)	
Organ	izer: Wei Chen (University of Pitts	burgh School of Medicine)	
13:30 – 13:	55 Heng Huang (University of I Deep Learning for Biomedical	[⊃] ittsburgh) Data Analysis	
13:55 – 14:	20 Yanming Li (University of M XG Boosting Variable Selection Time Varying Imaging Predic	ichigan) on for a Discrete Time Cure Rate Survival I tors	Model with High-Dimensional
14:20 - 14:	45 Ying Ding (University of Pitt A Copula-Based Semiparament tion (AMD) using GWAS Dat	tsburgh) tric Model for Progression Prediction of Ag ta	e-Related Macular Degenera-
Thursday	May 30, 13:30 – 15:00	Invited Session 11	BENEDUM G29
Founders S	Session on Current Topics in Lif	etime Data Analysis	

Chair and Organizer: Richard Cook (University of Waterloo, Canada)

- 13:30 13:55 Mei-Ling Ting Lee (University of Maryland, College Park) Distribution-free Threshold Regression for Longitudinal Time-to-event Analysis
 13:55 – 14:20 Jack Kalbfleisch (University of Michigan) Direct and Indirect Standardized Mortality Ratios Based on the Cox Model
- 14:20 14:45 **Mei-Cheng Wang** (Johns Hopkins University) Complexity in Simple Cross-Sectional Data with Binary Disease Outcome

Thursday May 30, 13:30 – 15:00

Invited Session 12

BENEDUM G30

Causal Inference in Life History Analysis

Chair and Organizer: Jon Michael Gran (University of Oslo, Norway)

13:30 – 13:55	Kjetil Røysland (University of Oslo, Norway) Causal Interpretation in Survival Analysis
13:55 – 14:20	Mats Julius Stensrud (University of Oslo, Norway) New Estimands for Causal Inference in the Presence of Competing Risks
14:20 - 14:45	Ruth Keogh (London School of Hygiene and Tropical Medicine) The sequential trials approach for estimating effects of treatment on survival using longitudinal observational data

ThursdayMay 30, 13:30 - 15:00Invited Session 13BENEDUM 158

Joint Models for Time-to-Event and Multiple Longitudinal Data or High Dimensional Data

Chair: Virginie Rondeau (Université de Bordeaux, France)

Organizer: Hélène Jacqmin-Gadda (Université de Bordeaux, France)

13:30 - 13:55	Sheng Luo (Duke University)
	Dynamic Prediction of Alzheimer's Disease Progression Using Features of Multiple Longitudinal Out-
	comes and Time-to-Event Data
13:55 – 14:20	Cecile Proust-Lima (Université de Bordeaux, France)
	Joint Modelling of Multiple Latent Processes and Clinical Endpoints: Application in Alzheimers Disease
14:20 - 14:45	Dimitris Rizopoulos (Erasmus University Medical Center, The Netherlands)

Invited Session 14

Using Joint Models for Personalized Optimal Scheduling of Invasive Procedures

Thursday May 30, 13:30 – 15:00

Survival Analysis with Error-Prone or Missing Covariate Measurements

Chair: Malka Gorfine (Tel Aviv University, Israel)

Organizer: Daniel Nevo (Tel Aviv University, Israel)

- 13:30 13:55 Daniel Nevo (Tel Aviv University, Israel)
 A Novel Calibration Framework for Survival Analysis When a Binary Covariate is Measured at Sparse Time Points
- 13:55 14:20 Xiao Song (University of Georgia) Partially Time-Varying Coefficient Proportional Hazards Models with Error-Prone Time-Dependent Covariates -an Application to the AIDS Clinical Trial Group 175 Data
- 14:20 14:45 **Molin Wang** (Harvard University) Semi parametric Methods for Exposure Misclassification in Propensity Score-Based Time-to-Event Data Analysis

Discussant: Malka Gorfine (Tel Aviv University, Israel)

Thursday May 30, 13:30 – 15:00

Novel Methods for the Analysis of Recurrent Event Data

Chair and Organizer: Douglas Schaubel (University of Michigan)

- 13:30 13:55 **Jianwen Cai** (University of North Carolina at Chapel Hill) Analysis of Recurrent Events Data from Case-Cohort Studies
- 13:55 14:20 **Chiung-Yu Huang** (University of California, San Francisco) Recurrent Events Analysis with Data Collected at Informative Clinical Visits
- 14:20 14:45 **Sehee Kim** (University of Michigan) Risk Discrimination Indexes for Recurrent Event Models : An Application to Identify Modifiable Risk Factors of Hospitalizations Among Dialysis Patients

BENEDUM G36

Invited Session 15

BENEDUM G27

Thursday May 30, 13:30 – 15:00

New Methods for Complex Censored Data

Chair: Chien-Yu Peng (Academia Sinica, Taiwan) Organizer: Mei-Ling Ting Lee (University of Maryland, College Park)

- 13:30 13:50 **Chung-Chou H. (Joyce) Chang** (University of Pittsburgh) Modeling Exposure-Time-Response Association in the Presence of Competing Risks
- 13:50 14:10 **Catherine Huber** (University of Paris, France) Efficient Semiparametric Maximum Likelihood Estimation (SPMLE) Under Interval Censoring and Truncation

Invited Session 16

- 14:10 14:30 **Ying Qing Chen** (University of Washington) Data Enriched Regression for Censored Time-to-Event
- 14:30 14:50 **Jonathan Race** (Ohio State University) Semiparametric Bayes Testing of Ordinal Effects on Survival

Thursday May 30, 15:00 – 15:30

Thursday May 30, 15:00 – 15:30

15:00 – 15:30 Refreshment Break

Thursday May 30, 15:30 – 17:00

Thursday May 30, 15:30 – 17:00 Invited Session 17 **BENEDUM G29** Advanced Statistical Methods for Time to Event Data in Complex Observational Studies Chair and Organizer: Jianwen Cai (University of North Carolina at Chapel Hill) 15:30 - 15:50Haibo Zhou (University of North Carolina at Chapel Hill) Secondary Analysis in Outcome Dependent Sampling Studies 15:50 - 16:10Yu Shen (University of Texas MD Anderson Cancer Center) Density Ratio Model for Analyzing Length-Biased Data **Qingning Zhou** (University of North Carolina at Charlotte) 16:10 - 16:30Two-stage Outcome-dependent Sampling with Interval-censored Failure Time Data 16:30 - 16:50**Donglin Zeng** (University of North Carolina at Chapel Hill)

Semiparametric Regression Analysis of Multiple Right- and Interval-Censored Events

Thursday May 30, 15:30 – 17:00

Recent Advances in Complex Bivariate Time-to-Event Data Modeling and Analysis

Chair and Organizer: Ying Ding (University of Pittsburgh)

- 15:30 15:55 **Yu Cheng** (University of Pittsburgh) Quantile Association Model for Bivariate Survival Data
- 15:55 16:20 **Joanna Shih** (NIH/NCI) Survival Analysis of Error-prone Self-report Time-to-event Outcome with an Internal Validation Sample Identified from a Mover-Stayer Model
- 16:20 16:45 **Jong Hyeon Jeong** (University of Pittsburgh) A Clustered Win Ratio for Semi-Competing Risks Data

BENEDUM G28

BENEDUM Hall Lobby

Invited Session 18

BENEDUM G30

Discussant: Richard Cook (University of Waterloo, Canada)

Thursday N	Nay 30, 15:30 – 17:00	Invited Session 19	BENEDUM 158
Complex Da	ta Analysis in the Biomedical	and Health Research	
Chair: \ Organiz	Wei Chen (University of Pittsburg er: Ruzong Fan (Georgetown Un	gh School of Medicine) iversity Medical Center)	
15:30 - 15:50	Christopher I. Amos (Baylo Understanding Complex Etiol	r College of Medicine) ogy of Autoimmune Diseases by Applicatior	n of Machine Learning Tools
15:50 - 16:10	Jason H. Moore (University Accessible Artificial Intelligen	of Pennsylvania) ce for Data Science	
16:10 - 16:30	 Saonli Basu (University Of N A Robust and Unified Frame mating Equations 	Лinnesota) work for Estimating Heritability in Twin St	udies using Generalized Esti-
16:30 - 16:50	Ruzong Fan (Georgetown Un Mixed Models for Gene-based	niversity Medical Center) I Association Analysis of Complex Traits	
Thursday N	Nay 30, 15:30 – 17:00	Invited Session 20	BENEDUM G36
Recent Adv	ances in Survival and Recurrer	ıt Event Data Analysis	
Chair ai	nd Organizer: Chiung-Yu Huang	(University of California, San Francisco)	
15:30 - 15:50	 Lili Wang (University of Mic Penalized Survival Models for 	higan) [,] the Analysis of Alternating Recurrent Ever	nt Data
15:50 - 16:10	Yifei Sun (Columbia Univers Survival Forest for Dynamic F	ity Mailman School of Public Health) Risk Prediction	
16:10 - 16:30	Zhezhen Jin (Columbia Univ On Time to Event Analysis for	versity Mailman School of Public Health) or Transplant Data	
16:30 - 16:50) Mi-Ok Kim (University of Ca Extended Likelihood Methods	alifornia, San Francisco) 5 for Censored Data	
Thursday N	Nay 30, 15:30 – 17:00	Invited Session 21	BENEDUM G27
Survival Ana	alysis Methods for Alzheimer's	Disease	
Chair a	nd Organizer: Lei Liu (Washingto	on University in St. Louis)	
15:30 - 15:50	Sharon X. Xie (University of Adjusting for Covariate Mea Applications to Alzheimer's D	f Pennsylvania) surement Error in Failure Time Analysis I Jisease Biomarker Research	Jnder Competing Risks with
15:50 - 16:10	Guoqiao Wang (Washingtor Survival Analysis for Alzheime	ı University in St. Louis) er Disease Without "Known Survival" Time	
16:10 - 16:30 16:30 - 16:50	 Dandan Liu (Vanderbilt Univ Partly Conditional Modeling f Lei Liu (Washington Univers Variable Selection in Joint Fr. 	versity) for Alzheimer's Disease ity in St. Louis) ailty Model of Recurrent and Terminal Ever	nts

Mediation Analysis for High-Dimensional Data

Chair and Organizer: Ruth Pfeiffer (NIH/NCI)

- 15:30 15:50 **Joshua Millstein** (University of Southern California) A Modification to the Causal Inference Test (CIT) Addresses Challenges of Mediation Analysis in Genomic Settings
- 15:50 16:10 Michael B. Sohn (University of Rochester) A Compositional Mediation Model for Microbiome Data
- 16:10 16:30 Andriy Derkach (NIH/NCI) High Dimensional Mediation Analysis
- 16:30 16:50 Xi Luo (Brown University) Mediation Analysis for Large and Multilevel Data

Thursday May 30, 15:30 – 17:00 Invited Session 23

Innovative Applications of Joint Modeling

Chair and Organizer: Rajeshwari Sundaram (NIH/NICHD)

Qing Pan (George Washington University) Constrained Maximum Entropy Models to Select Genotype Interactions Associated with Survival Outcomes
Sedigheh Mirzaei (St. Jude Children's Research Hospital) Statistical Modeling of Discrete Survival Time in Presence of Recall Bias: A Joint Modeling Approach
Angelo Elmi (George Washington University) Approximate Imputation Based Methods for Joint Modeling of Longitudinal and Time-to-Event Data
Li Cheung (NIH/DCEG) Mixture Models for Undiagnosed Prevalent Disease and Interval Censored Incident Disease: Applications to a Cohort Assembled from Electronic Health Records

Invited Session 24

Thursday May 30, 15:30 – 17:00

Applied Stochastic Models for Time-to-Event Data

Chair: Henry W. Block (University of Pittsburgh) Organizer: Mei-Ling Ting Lee (University of Maryland, College Park)

- 15:30 15:55 **Satish Iyengar** (University of Pittsburgh) Diffusion Models for Time-to-Event Data
- 15:55 16:20 **Chien-Yu Peng** (Academia Sinica, Taiwan) Robust Processes in Degradation Analysis
- 16:20 16:45 **Tianzhou Ma** (University of Maryland, College Park) Variable Selection in Threshold Regression Model with Application to HIV Drug Adherence Data

Thursday May 30, 18:00 – 20:30

Thursday May 30, 18:00 – 20:30

18:00 – 20:30 Banquet

Wyndham Hotel

BENEDUM G37

BENEDUM G28

Friday May 31

	31, 8:00 - 9:00	Keynote Presentation	BENEDUM 157
Keynote Pres	sentation: Ross Prentice		
Chair: Ji	anwen Cai (University of Nor	th Carolina at Chapel Hill)	
8:00 – 9:00	Ross Prentice (Fred Hutchin Regression Models and Multi	nson Cancer Research Center/University of W variate Life Tables	ashington)
Friday May 3	31, 9:00 – 10:00	Keynote Presentation	BENEDUM 157
Keynote Pres	sentation: Danyu Lin		
Chair: Ji	anwen Cai (University of Nor	th Carolina at Chapel Hill)	
9:00 - 10:00	Danyu Lin (University of N Semiparametric Regression)	lorth Carolina at Chapel Hill) Analysis of Interval-Censored Data	
Friday Ma	y 31, 10:00 - 10:30		
Friday May	31, 10:00–10:30		BENEDUM Hall Lobby
Friday May 3	31, 10:00–10:30 Refreshment Break		BENEDUM Hall Lobby
Friday May 3 10:00 – 10:30 Friday Ma	31, 10:00–10:30 Refreshment Break y 31, 10:30 – 12:00		BENEDUM Hall Lobby
Friday May 3 10:00 – 10:30 Friday Ma Friday May 3	 31, 10:00–10:30 Refreshment Break y 31, 10:30 – 12:00 31, 10:30 – 12:00 	Invited Session 25	BENEDUM Hall Lobby BENEDUM G30
Friday May 3 10:00 – 10:30 Friday Ma Friday May 3 Innovative Aj	 31, 10:00–10:30 Refreshment Break y 31, 10:30 – 12:00 31, 10:30 – 12:00 pproaches for Studying the	Invited Session 25 e Effects of Chemical Mixtures on Disease	BENEDUM Hall Lobby BENEDUM G30 Throughout the Life-span
Friday May 3 10:00 – 10:30 Friday Ma Friday May 3 Innovative A Chair and	 31, 10:00–10:30 Refreshment Break y 31, 10:30 – 12:00 31, 10:30 – 12:00 pproaches for Studying the d Organizer: Paul Albert (NII) 	Invited Session 25 e Effects of Chemical Mixtures on Disease H/NCI)	BENEDUM Hall Lobby BENEDUM G30 Throughout the Life-span
Friday May 3 10:00 – 10:30 Friday Ma Friday May 3 Innovative A Chair and 10:30 – 10:55	 31, 10:00–10:30 Refreshment Break y 31, 10:30 – 12:00 31, 10:30 – 12:00 pproaches for Studying the d Organizer: Paul Albert (NII Shelley Liu (Icahn School Bayesian Methods for Anal	Invited Session 25 e Effects of Chemical Mixtures on Disease H/NCI) of Medicine at Mount Sinai) lyzing Chemical Mixtures	BENEDUM Hall Lobby BENEDUM G30 Throughout the Life-span
Friday May 3 10:00 – 10:30 Friday Ma Friday May 3 Innovative A Chair and 10:30 – 10:55 10:55 – 11:20	 31, 10:00–10:30 Refreshment Break y 31, 10:30 – 12:00 31, 10:30 – 12:00 pproaches for Studying the d Organizer: Paul Albert (NII Shelley Liu (Icahn School Bayesian Methods for Anal David Wheeler (Virginia Models for Environmental Models for Environmental	Invited Session 25 e Effects of Chemical Mixtures on Disease H/NCI) of Medicine at Mount Sinai) lyzing Chemical Mixtures Commonwealth University) Chemical Mixtures and Childhood Leukemia F	BENEDUM Hall Lobby BENEDUM G30 Throughout the Life-span
Friday May 3 10:00 – 10:30 Friday Ma Friday May 3 Innovative A Chair and 10:30 – 10:55 10:55 – 11:20 11:20 – 11:45	 31, 10:00–10:30 Refreshment Break y 31, 10:30 – 12:00 31, 10:30 – 12:00 pproaches for Studying the d Organizer: Paul Albert (NII Shelley Liu (Icahn School Bayesian Methods for Anal David Wheeler (Virginia Models for Environmental Zhen Chen (NIH/NICHD) A Bayesian Multi-dimensio	Invited Session 25 e Effects of Chemical Mixtures on Disease H/NCI) of Medicine at Mount Sinai) lyzing Chemical Mixtures Commonwealth University) Chemical Mixtures and Childhood Leukemia F) mal Couple-based Latent Risk Model with an	BENEDUM Hall Lobby BENEDUM G30 Throughout the Life-span Risk

Friday May 31, 10:30 – 12:00

Invited Session 26

BENEDUM 158

Chair and Organizer: Rebecca A. Betensky (New York University)

10:30 - 10:55	Lior Rennert (Clemson University)
	Cox Regression Model under Dependent Truncation
10:55 - 11:20	Jing Qian (University of Massachusetts, Amherst)

Estimation under Covariate-Induced Dependent Truncation with Inverse Probability of Weighting

Bella Vakulenko-Lagun (Harvard University) 11:20 - 11:45Nonidentifiability in the Presence of Factorization for Truncated Data

Discussant: Rebecca A. Betensky (New York University)

Friday	May 31, 10:30 – 12:00	Invited Session 27	BENEDUM G29

Methods for Lifetime Data Processes Under Intermittent Observation

Chair and Organizer: Richard Cook (University of Waterloo, Canada)

- 10:30 10:55Jerald Lawless (University of Waterloo, Canada) Independence Conditions and Life History Analysis with Intermittent Observation of Individuals 10:55 - 11:20 **Pamela Shaw** (University of Pennsylvania) Assessing efficacy for an interval-censored bivariate failure time outcome when one event is a surrogate
- Andrew Titman (Lancaster University, UK) 11:20 - 11:45 A Joint Models for Multi-state Models with Informative Observation Processes

Friday May 31, 10:30 – 12:00

Cross-Sectional Analysis of Life History Data

Chair and Organizer: X. Joan Hu (Simon Fraser University, Canada)

- Leilei Zeng (University of Waterloo, Canada) 10:30 - 10:55Multistate Analysis from Cross-Sectional and Auxiliary Samples
- 10:55 11:20Niels Keiding (University of Copenhagen, Denmark) Possible Application of Current Duration Analysis to Estimate Time-to-Pregnancy Distributions from Demographic Survey Data
- 11:20 11:45 **Cuiling Wang** (Albert Einstein College of Medicine) Evaluate Cross-Sectional Association and Diagnostic Accuracy for Disease using Longitudinal Markers with Missing Data

Friday May 31, 10:30 - 12:00

Novel Approaches for Time-to-Event Data

Chair: Guoqing Diao (George Mason University) Organizer: Aiyi Liu (NIH/NICHD)

- 10:30 10:55 Wei Qian (University of Delaware) Double-slicing Assisted Sufficient Dimension Reduction for High Dimensional Censored Data 10:55 - 11:20**Rajeshwari Sundaram** (NIH/NICHD) Analyzing Recurrent Events in Presence of Recall Error : An Application to Time-to-repeat Hospitalization
- 11:20 11:45 **Antai Wang** (New Jersey Institute of Technology) Analysis of Semi-competing Risks Data using Archimedean Copula Models

BENEDUM 226

Invited Session 29

Invited Session 28

BENEDUM 227

Friday	May 31	, 10:30 - 12:00	Invited Session 30	BENEDUM G27
Multipl	e Testir	ıg, Simultaneous and Jo	int Modeling, and Calibration	
Ch	air: Shiv	wen Shen (New York Life Edsel A. Peña (University	Insurance)	
10:30 -	10:55	Yen-Yi Ho (University of P value Adjustment Proc	f South Carolina) edure Using Empirical Weights	
10:55 –	11:20	Piaomu Liu (Bentley Un Joint Dynamic Modeling	iversity) of a Longitudinal Marker, Recurrent Competing	g Risks and a Terminal Event
11:20 -	11:45	Beidi Qiang (Southern II Estimating Concentration Data	linois University - Edwardsville) Response Function and Change-Point using	Time-Course and Calibration
Friday	May 31	, 10:30 – 12:00	Invited Session 31	BENEDUM G28

Genomic Applications with Survival Outcomes

Chair: Tianzhou Ma (University of Maryland, College Park)

Organizer: Mei-Ling Ting Lee (University of Maryland, College Park)

- 10:30 10:55 **George Tseng** (University of Pittsburgh) Heterogeneity and Robust Machine Learning in Transcriptomic Analysis of Multiple Studies
- 10:55 11:20 Philippe Broët (INSERM, France)
 Predicting the Occurrence of an Adverse Event using Bagged Survival Trees Built on Random Structured Subsets: Application in Clinical Immunology
- 11:20 11:45 **Chien-Wei Lin** (Medical College of Wisconsin) RNASeqDesign: A Framework for RNA-Seq Genome-wide Power Calculation and Study Design Issues

Invited Session 32

Invited Session 33

Friday May 31, 10:30 – 12:00

New Methods for Risk Prediction and Precision Medicine

Chair and Organizer: Mei-Cheng Wang (Johns Hopkins University)

- 10:30 10:55 Yanxun Xu (Johns Hopkins University) Bayesian estimation of Individualized Treatment-Response Curves in Electronic Medical Record Data
- 10:55 11:20Jon Arni Steingrimsson (Brown University)
Deep Learning with Time-to-event Outcomes
- 11:20 11:45 **Xingqiu Zhao** (Hong Kong Polytechnic University, Hong Kong) Subgroup Analysis in the Cox Model

Friday May 31, 10:30 – 12:00

Causal Inference for Survival Data

Chair and Organizer: Ronghui (Lily) Xu (University of California, San Diego)

- 10:30 10:55 **Eric J. Tchetgen Tchetgen** (University of Pennsylvania) Mendelian Randomization for Failure Time Outcome with Invalid Instrumental Variables
- 10:55 11:20 **Jessica Young** (Harvard Pilgrim Health Care Institute) A Causal Framework for Classical Statistical Estimands in Failure Time Setting with Competing Events
- 11:20 11:45 **Jelena Bradic** (University of California, San Diego) Inference on Treatment Effects in High-dimensional Survival Models

BENEDUM G31

BENEDUM G36

Friday May 31, 12:00 – 13:30

Friday May 31, 12:00 - 13:30

12:00 – 13:30 Lunch Break

Friday May 31, 13:30 – 15:00

Friday	May 31	l, 13:30 – 15:00	Invited Session 34	BENEDUM 226
Session	for Stu	ident Paper Award Wir	iners (1)	
Ch	air: Guo	oqing Diao (George Maso	n University)	
Or	ganizer:	Guoqing Diao (George N	1ason University)	
13:30 -	13:50	Ting Ye (University of N Robust Tests for Treatm	Visconsin-Madison) ent Effect in Survival Analysis under Covariate-/	Adaptive Randomization
13:50 –	14:10	Gabrielle Simoneau (M Estimating Optimal Dyn	cGill University, Canada) amic Treatment Regimes with Survival Outcom	es
14:10 -	14:30	Rui Chen (University of Tailored Optimal Post-T	Wisconsin-Madison) reatment Surveillance for Cancer Recurrence	
14:30 -	14:50	Nicole Butera (Universi Estimating Biomarker Ch	ty of North Carolina at Chapel Hill) nange by Adjusting for Informative Time to Trea	atment Initiation
Friday	May 31	l, 13:30 – 15:00	Invited Session 35	BENEDUM G30
Causal	Inferen	ce with Time-to-Event	Data	

Chair and Organizer: Lu Mao (University of Wisconsin-Madison)

- 13:30 13:55 **Mark van der Laan** (University of California, Berkeley) Target MLE of Causal Effects of Interventions on Survival Outcomes
- 13:55 14:20 **Yifan Cui** (University of Pennsylvania) Instrumental Variable Estimation of Marginal Structural Cox Model for Time-Varying Exposure
- 14:20 14:45 **Constantine Frangakis** (Johns Hopkins University) Deductive Semiparametric Estimation in Double-Sampling Designs with Application to Estimating Mortality in PEPFAR

Discussant: Lu Mao (University of Wisconsin-Madison)

Friday May 31, 13:30 – 15:00 Invited Session 36

BENEDUM G31

Multistate Models as A Framework for Life History Analysis

Chair and Organizer: Hein Putter (Leiden University Medical Centre, The Netherlands)

- 13:30 13:50 Hans C. van Houwelingen (Leiden University Medical Center, The Netherlands) An Alternative for the Fine-Gray Approach to Competing Risk Modelling: a Bridge Between Multi-State and Subdistribution Hazard
 13:50 – 14:10 Ronald Geskus (University of Oxford at Ho Chi Minh City, Vietnam)
- A Bayesian Analysis of the Natural History of HIV-2 Infection using a Hidden Markov Cure Model
- 14:10 14:30 **Jeremy Taylor** (University of Michigan) Handling Missing Covariate Data in Multistate Models
- 14:30 14:50 **Jon Michael Gran** (University of Oslo, Norway) Estimating Causal Effects of Time-varying Treatments in Multi-State Models - An Application to Registry Data on Sick Leave and Disability

Friday May 31, 13:30 – 15:00

Models and Applications with Recurrent Events

Chair and Organizer: Thomas Scheike (University of Copenhagen, Denmark)

- 13:30 13:55 **Virginie Rondeau** (Universitè de Bordeaux, France) Multivariate Joint Frailty Model for the Analysis of Nonlinear Tumor Kinetics with Recurrent Progressions of Nontarget Progression and Dynamic Predictions of Death
- Per Kragh Andersen (University of Copenhagen, Denmark) 13:55 - 14:20Marginal Regression Models for Recurrent Events with Competing Risks using Pseudo-Obervations
- 14:20 14:45**Frank Eriksson** (University of Copenhagen, Denmark) The Mean, Variance, and Correlation for Bivariate Recurrent Events Data with a Terminal Event

Friday May 31, 13:30 - 15:00 **Invited Session 38 BENEDUM 158**

Challenges and Advances of Research in Health Service Studies

Chair: Liang Zhu (University of Texas Health Science Center at Houston)

- Organizer: Yu Shen (University of Texas MD Anderson Cancer Center)
- 13:30 13:55 Jane Lange (Fred Hutchinson Cancer Research Center) Estimating and Comparing Cancer Progression Risks Under Varying Surveillance Protocols : Moving Beyond the "Tower of Babel"
- 13:55 14:20 Nabihah Tayob (University of Texas MD Anderson Cancer Center) Longitudinal Biomarker Screening Algorithms: Guidelines for When They are Most Useful
- Liang Li (University of Texas MD Anderson Cancer Center) 14:20 - 14:45 Modeling the Longitudinal Trajectory of Medical Cost for Cancer Care

Friday May 31, 13:30 - 15:00

Innovative Methods for Assessing Diagnostic Accuracy and Prediction Accuracy

Chair: Sedigheh Mirzaei (St Jude Children's Research Hospital)

Organizer: Rajeshwari Sundaram (NIH/NICHD)

- 13:30 13:55 Paramita Saha-Chaudhuri (McGill University, Canada) Monitoring with Repeatedly Measured Marker: Assessing Incremental Value of Additional Measurements
- Aasthaa Bansal (University of Washington) 13:55 - 14:20 Evaluating the Time-varying Prediction Accuracy of Survival Models Used in Dynamic Decision-making
- 14:20 14:45 Paramita Saha-Chaudhuri (McGill University, Canada) Mean Risk Stratification and Number Needed to Test: A New Approach to Quantifying Risk Stratification for Comparing the Usefulness of Diagnostic Tests

Friday May 31, 13:30 - 15:00

Invited Session 40

BENEDUM G28

BENEDUM G27

Extensions to Joint Longitudinal-Survival Modelling

Chair and Organizer: Andrew Titman (Lancaster University, UK)

BENEDUM G29

Invited Session 39

Invited Session 37

13:30 – 13:55	Li Su (University of Cambridge, UK) Accommodating Informative Dropout and Death: A Joint Modelling Approach for Longitudinal and Semicompeting Risks Data
13:55 – 14:20	Zhigang Li (University of Florida) Joint Modeling Quality of Life and Survival in Palliative Care Research
14:20 – 14:45	Bin Nan (University of California, Irvine) Conditional Modeling of Longitudinal Data with Terminal Event

Friday May 31, 13:30 – 15:00 **Invited Session 41 BENEDUM G36**

Emerging Issues and Methods on Censored Data

Chair and Organizer: Grace Y. Yi (University of Waterloo, Canada)

- 13:30 13:55 Masoud Asgharian (McGill University, Canada) Prevalent Cohort Studies: Length-Biased Sampling with Right Censoring
- 13:55 14:20 **Douglas Schaubel** (University of Michigan) Semiparametric Regression Methods for Temporal Processes Subject to Multiple Sources of Censoring
- 14:20 14:45 **Gang Li** (University of California, Los Angeles) Ultrahigh Dimensional Screening for Survival Data

Discussant: Ronghui (Lily) Xu (University of California, San Diego)

Invited Session 42 Friday May 31, 13:30 - 15:00

Intermittent Observation of Life History Processes

Chair and Organizer: Leilei Zeng (University of Waterloo, Canada)

- 13:30 13:55 **X. Joan Hu** (Simon Fraser University, Canada) Statistical Issues in Administrative Data Analysis
- 13:55 14:20 Yueh-Ying Han (University of Pittsburgh) Utilizing National Health Survey Data for Surviving Analysis
- 14:20 14:45 Eleanor Pullenayegum (Hospital of Sick Children, Canada) The Role of Recurrent Event Models in the Analysis of Longitudinal Data Subject to Irregular Observation: Current Practice and Future Directions

Friday May 31, 15:00 – 15:30

Friday May 31, 15:00 – 15:30

15:00 – 15:30 Refreshment Break

Friday May 31, 15:30 – 17:00

Friday May 31, 15:30 - 17:00 **Multi-State Models in Practice**

Chair: Richard Cook (University of Waterloo, Canada) Organizer: Mouna Akacha (Novartis, Switzerland)

BENEDUM Hall Lobby

BENEDUM 227

Invited Session 43

BENEDUM 158

15:30 - 15:50	David James (Novartis, US) Multistate Modeling and Simulation of Patient Trajectories After Allogeneic Hematopoietic Stem Cell Transplantation to Inform Drug Development
15:50 - 16:10	Ulrich Beyer (Roche, Switzerland) A Multistate Model for Early Decision Making in Oncology
16:10 - 16:30	Jan Feifel (Ulm University, Germany) Utilization of Multistate Models and Subcohorting to Analyze Rare Exposures
16:30 - 16:50	Terry Therneau (Mayo Clinic) Practical Multi-State Models

Friday May 31, 15:30 – 17:00 Invited Session 44

Session for Student Paper Award Winners (2)

Chair: Guoqing Diao (George Mason University) Organizer: Guoqing Diao (George Mason University)

15:30 – 15:50 **Bo Wei** (Emory University) Generalized Accelerated Recurrence Time Model in the Presence of a Dependent Terminal Event

- 15:50 16:10 **Yi Xiong** (Simon Fraser University, Canada) Estimating Duration Distribution from Data with Missing Time Origin
- 16:10 16:30 Yue Wei (University of Pittsburgh) Gene-based Association Analysis for Bivariate Time-to-event Data through Functional Regression with Copula Models

Invited Session 45

Invited Session 46

Friday May 31, 15:30 – 17:00

Recent Developments in Statistical Methods on Semi-Competing Risks Data

Chair and Organizer: Jong Hyeon Jeong (University of Pittsburgh)

15:30 – 15:55 David Oakes (University of Rochester) Nonparametric Estimation of the Curtailed Win-Ratio
15:55 – 16:20 Lu Mao (University of Wisconsin-Madison) On the Win-Loss Processes of Composite Endpoints
16:20 – 16:45 II-Do Ha (Pukyong National University, South Korea) H-likelihood Approaches for Frailty Models with Semi-competing Risks Data

Friday May 31, 15:30 - 17:00

Survival Analysis with Missing or Mismeasured Data

Chair and Organizer: Bin Nan (University of Michigan)

- 15:30 15:55 Ronghui (Lily) Xu (University of California, San Diego) Two-Stage Residual Inclusion for Survival Data and Competing Risks - An Instrumental Variable Approach for Binary Treatment with Application to SEER-Medicare Linked Data
 15:55 – 16:20 Grace Y. Yi (University of Waterloo, Canada)
- Analysis of Error-Prone Survival Data Under Additive Hazards Models
- 16:20 16:45 **Yichuan Zhao** (Georgia State University) Rank-based Estimating Equation with Non-ignorable Missing Responses

BENEDUM 227

BENEDUM G31

BENEDUM G30

Friday	May 31	l, 15:30 – 17:00	Invited Session 47	BENEDUM G27
Recent	Advand	ces in the Analysis of	Complex Lifetime Data Involving Recurrent Events	
Cł	nair and	Organizer: Hua Shen (U	Jniversity of Calgary, Canada)	
15:30 –	15:55	Ming Wang (Pennsylv A Time-varying Joint F Application to the Carc	rania State University) Frailty-copula Model for Analyzing Recurrent Events and a liovascular Health Study	Terminal Event: An
15:55 –	16:20	Kaida Cai (University Bi-level Variable Selecti	of Calgary, Canada) ion for Multivariate Failure Time Data with Observed Het	erogeneity
16:20 –	16:45	Shu Jiang (Harvard U Finite Mixture Models	niversity) for Multistate Processes under Panel Observation	
Friday	May 3	l, 15:30 – 17:00	Invited Session 48	BENEDUM G28
Person	alized T	reatment Selection wi	ith Censored Survival Outcome	
Cł	nair and	Organizer: Xiao Song (I	University of Georgia)	
15:30 –	15:55	Abdus S. Wahed (Un Optimizing Dynamic Tr	iversity of Pittsburg) reatment Regimes Based on Quality-Adjusted Survival	
15:55 –	16:20	Yingqi Zhao (Fred Hu Constructing Stabilized	tchinson Cancer Research Center) Dynamic Treatment Regimes for Censored Data	
16:20 –	16:45	Lihui Zhao (Northwest Identifying Potential Re	tern University) esponses to a New Treatment with Time-to-event Endpoir	its
Friday	May 3	l, 15:30 – 17:00	Invited Session 49	BENEDUM G36
Recent	Develo	pment in the Analysis	of Recurrent Event Data	
Cł	nair: Qin	gning Zhou (University	of North Carolina at Charlotte)	
0	rganizer:	Jianguo (Tony) Sun (U	Iniversity of Missouri)	
15:30 –	15:55	Yang Li (University of A General Additive-Mu	North Carolina at Charlotte) Itiplicative Mean Model for Panel Count Data Analysis	
15:55 –	16:20	Liang Zhu (University Statistical Analysis on I	of Texas Health Science Center at Houston) Mixed Recurrent Event Data with Clusters	
16:20 –	16:45	Shijun Zhu (University A Joint Model of Longi	v of Maryland, Baltimore Country) itudinal Biomarkers and Recurrent Events and it's Applica	tion
Friday	May 3	l, 15:30 – 17:00	Invited Session 50	BENEDUM G29

Novel Application of Survival Models in Complex Biomedical Studies

Chair and Organizer: Donglin Zeng (University of North Carolina at Chapel Hill)

 15:30 – 15:55 Yuanjia Wang (Columbia University Mailman School of Public Health) Early Diagnosis of Neurological Disease Using Peak Degeneration Ages of Multiple Biomarkers
 15:55 – 16:20 Fei Gao (University of Washington)

Semiparametric Regression Analysis of Length-Biased Interval-Censored Data

 16:20 – 16:45 Qingxia Chen (Vanderbilt University)
 Treatment Effect Estimate and Model Diagnostics with Two-way Time-Varying Treatment Switching: An Application to a Head and Neck Study

Two-Phase Studies for Lifetime Data

Ørnulf Borgan (University of Oslo Norway) Sven Ove Samuelsen (University of Oslo, Norway)

Overview

In cohort studies, regression methods are commonly applied to assess the influence of risk factors and other covariates on mortality or morbidity; in particular Cox-regression is much used. Estimation in Coxs model is based on a partial likelihood that at each observed death or disease occurrence ("failure") compares the covariate values of the failing individual to those of all individuals at risk. Thus Cox regression requires collection of covariate information for all individuals in the cohort, even when only a small fraction of them actually get diseased or die. This may be very expensive, or even logistically impossible. Further, when covariate measurements are based on biological material stored in biobanks, it will imply a waste of valuable material that one may want to save for future studies. Cohort sampling designs, where covariate information is collected for all failing individuals ("cases"), but only for a sample of the individuals who do not fail ("controls"), then offer useful alternatives that may save biological material and drastically reduce the workload of data collection and error checking. Such cohort sampling designs may be considered as two-phase designs, where the cohort is the phase I sample (selected from a superpopulation) and the case-control sample is the phase II sample selected from the cohort.

There are two main types of two-phase designs for life time data: nested case-control and case-cohort designs, and the two types of designs differ in the way controls are selected. The course presents the two types of designs both in their original form and later extensions and describes how the statistical analysis of such two-phase studies may be performed. The focus is on estimation of relative risks using partial likelihoods and pseudo-likelihoods (or weighted likelihoods) that resemble the full cohort partial likelihood. Other topics like estimation of absolute risk and model checking will also be discussed, and methods that use all available data in the full cohort will be mentioned. There will be practical exercises in analyzing two-phase life time data, and the participants should bring their own laptop with R installed. Information on R packages that are needed will be given closer to the course.

Aims

1. Introduce the most common two-phase designs for life time data: nested case-control and case-cohort.

- 2. Discuss classical statistical methods for estimating relative risks for two-phase life time data, and give an outline of their theoretical properties.
- 3. Discuss methods for absolute risk estimation and model assessment.
- 4. Describe two-phase methods that use all available data from the full cohort.
- 5. Illustrate how to carry out statistical analyses of two-phase life time data using R.

Learning Outcomes

At the end of the day participants should:

- 1. Know the characteristics of the two common types of two-phase designs for life time data and understand the pros and cons of the designs.
- 2. Know how to estimate relative and absolute risks from nested case-control and case-cohort data.
- 3. Have some knowledge of methods that make use of data that are available for the full cohort.
- 4. Have some experience in analyzing nested case-control and case-cohort data using R.

Topics Covered

The material will be presented in a lecture format, where the theory and methods will be motivated and illustrated by examples from health research. In addition, the participants will get hands-on experience with the methods from practical exercises using R. Topics covered include:

1. Summary of methods for analyzing cohort life time data.

- 2. Nested case-control designs, including counter-matched sampling of the controls.
- 3. Case-cohort designs, including stratified sampling of the subcohort.
- 4. Classical methods for estimating relative and absolute risk from nested case-control and case-cohort data.
- 5. Analysis of general models for nested case-control and case-cohort data using inverse probability weighting.
- 6. Calibration of inverse probability weights for case-cohort data.
- 7. Methods for two-phase data that use all available cohort information (multiple imputation and maximum likelihood).
- 8. Practical examples and exercises using R.

Learning Strategy

The material will be presented using slides, class discussion, and practical exercises using R. Attendees will be given a booklet containing the slides, which will contain clear descriptions of the methodology, of applications, and of how to implement analyses in R.

Pre-requisites

The short course will be directed at statisticians in academia, government or industry interested in learning about two phase designs for life time data. It will be assumed that the participants are familiar with the basic concepts and methods in survival analysis and that they have some experience in using the R software.

Recommended reading

- Chapters 7 and 8 of Keogh & Cox: Case-Control Studies, Cambridge University Press, 2014.
- Part IV of *Handbook of Statistical Methods for Case-Control Studies*, eds Borgan, Breslow, Chatterjee, Gail, Scott & Wild, CRC Press, 2018.

About the Instructors



Ørnulf Borgan is professor of Statistics at the University of Oslo. His main research interest has been statistical methods for survival and event history data, including nested case-control and case-cohort designs. He is co-author of two books on the use of counting processes and martingales in survival and event history analysis, and he is one of the editors of the recent *Handbook of Statistical Methods for Case-Control Studies* (CRC Press, 2018). Borgan has been editor of the Scandinavian Journal of Statistics, and he is a Fellow of the American Statistical Association and member of the Norwegian Academy of Science and Letters.



Sven Ove Samuelsen is professor of Statistics at the University of Oslo. His main research interest has been statistical methods for survival and event history data, in particular case-cohort and nested case-control designs. He has been involved in planning and analyzing many case-control and other epidemiological studies. Samuelsen is on the editorial board of Lifetime Data Analysis.

Dynamic Prediction in Survival Analysis

Hein Putter (Leiden University Medical Center, The Netherlands)

Summary

The medical literature abounds with prediction models. They are statistical models based on patient- and disease characteristics, used to inform treatment decisions, to provide personalized risk estimates for the patient, and also to stratify patients in clinical trials. Important prognostic models include Adjuvant! Online in cancer and the Framingham risk score in cardiovascular disease. The vast majority of these models are focused on prognosis at one well-defined baseline moment, typically at diagnosis, shortly before treatment is initiated. It is at this time that the most important decisions on primary treatment are made. There is little doubt that the available prognostic models are important tools for the treating physician to guide treatment decisions at diagnosis. However, once primary treatment has been initiated, the prognosis of the patient will change over the course of time, as a result of the effect of treatment, possible treatment toxicity, and clinical events such as disease recurrence that may have occurred, and, very simply, because of the fact that the patient is still alive. As a result, these prediction models need to be "updated" to use the knowledge that has become available since baseline. Prediction models that incorporate this dynamic aspect are called dynamic prediction models, and they are the topic of this course.

This course will focus on methodology for dynamic prediction. The dynamic aspect of dynamic prediction involves using information on events and/or measurements up to the present, in order to "update" the prediction. It will be shown in this course how dynamic predictions may be obtained using the concept of landmarking and using multi-state models. Analyses will be illustrated using R, in particular the mstate and dynpred packages. Implementation of the methods in other statistical software packages like SAS, Stata and SPSS will be discussed.

Aims

- 1. Discuss situations where dynamic prediction is relevant;
- 2. Illustrate how the Cox model can be used to obtain dynamic predictions with time-fixed covariates;
- 3. Introduce multi-state models as an extension of survival analysis and competing risks;
- 4. Show how multi-state models can be used to obtain dynamic predictions;
- 5. Introduce landmarking as a way of dealing with time-dependent covariates;
- 6. Show how landmarking can be used to include time-dependent information in the dynamic predictions;
- 7. Discuss robustness properties;
- 8. Illustrate how to carry out the analyses discussed during the course using R.

Learning Outcomes

At the end of the course participants should:

- 1. Understand the connection between hazards and dynamic prediction probabilities;
- 2. Know how to obtain dynamic prediction probabilities from time-fixed Cox models;
- 3. Understand the difficulties of predicting with time-dependent covariates;
- 4. Be acquainted with concepts in multi-state models like transition intensities, transition probabilities, state occupation probabilities, the Markov assumption;
- 5. Understand the relation between transition intensities and transition probabilities, and be acquainted with the Aalen-Johansen estimator;
- 6. Understand how landmarking can be used for dynamic prediction.

Topics Covered

The course material will be presented in a lecture format, changing between theory and illustrations. Ample attention will be devoted to the practical implementation of the methods covered in the course, using R.

Topics covered include:

• Dynamic use of familiar survival analysis techniques

A short overview of survival analysis will be given, including the Cox model. The emphasis in this overview will be on how these familiar techniques can be used to obtain dynamic predictions. We will introduce conditional survival (the effect of being alive) and the fixed width failure function, and their relation to the familiar hazard function. Extensions to competing risks will briefly be mentioned.

• Time-dependent covariates and landmarking

We will then introduce time-dependent covariates and discuss techniques to handle them such as time-dependent Cox regression and landmarking. The differences between these approaches and the relative merits will be discussed.

• Multi-state models

A brief overview of multi-state models will be given, including how they can be used to obtain dynamic predictions. The overview includes discussion of concepts like transition intensities and transition probabilities, and ways of estimating transition intensities. The Aalen-Johansen estimator of the transition probabilities will be presented, and the assumptions needed for validity of the Aalen-Johansen estimator, in particular the Markov assumption will be discussed.

• Landmarking and dynamic prediction

Then we will show how landmarking can be used to include time-dependent information in the dynamic predictions. We will briefly discuss more traditional methods that can also be used for dynamic prediction, such as multi-state models. Advantages and disadvantages of different approaches will be discussed.

• Practical implementation

Methods discussed during the lectures will be illustrated using R, and in particular the mstate and dynpred packages. Data used is available from the presenter upon request.

Learning Strategy

The material will be presented using slides and through class discussion. Attendees will be given a booklet containing the slides , which will contain clear descriptions of the methodology, of applications, and of how to implement analyses in R.

Pre-requisites

This course is directed at statisticians or epidemiologists in academia, government or industry interested in dynamic prediction in survival analysis. Participants are expected to have a fair knowledge of the techniques from classical survival analysis.

About the Instructor



Hein Putter is Professor at the Leiden University Medical Center (Department of Biomedical Data Sciences). His research interests include competing risks and multi-state models, frailty models and dynamic prediction. He is co-author of the book "Dynamic Prediction in Clinical Survival Analysis", with Hans van Houwelingen.

Biased Sampling, Left Truncation and Survival Analysis

Jing Qin (NIH/NIAID)

Overview

Biased sampling occurs when a proper randomization cannot be achieved, the observed sample will not be representative of the population of interest. Biased sampling problems appear in many areas of research, including, Medicine, Epidemiology and Public Health, Social Sciences and Economics. Left truncation and length-biased data are clearly encountered in applications of renewal processes, etiologic studies, genome-wide linkage studies, epidemiologic cohort studies, cancer prevention trials, and studies of labor economy. In observational studies, a prevalent cohort design that draws samples from individuals with a condition or disease at the time of enrollment is generally more efficient and practical. The recruited patients who have already experienced an initiating event are followed prospectively for the failure event (e.g. disease progression or death) or are right censored. Under this sampling design, individuals with longer survival times measured from the onset of the disease are more likely to be included in the cohort, whereas those with shorter survival times are unconsciously excluded. Finding appropriate adjustments for the potential selection bias in analyzing length-biased data or more general biased sampling problems has been a long standing statistical problem.

This workshop discusses various methods to deal with biased sampling problems, exponential tilting models and left truncation and right censored data problems, including prole maximum likelihood method, conditioning likelihood method, composite partial likelihood method as well as general imputation methods.

Aims and Topics Covered

- 1. Discuss the general methods for handling biased sampling problems, including case and control problems, missing data and casual inference.
- 2. Derive the Cox partial likelihood from different angles, including rank likelihood method, prole maximum likelihood method, case and control conditional likelihood argument, and optimal estimating equation method.
- 3. Present the latest results on analyzing length biased survival time data, including Vardi's multiplicative censoring problem, and general imputation principle for missing survival data.
- 4. Discuss the well known pool adjacent violators algorithm and its combination with the EM algorithm together for estimating shape constrained inference, including estimation of monotonic decreasing density, cumulation hazard or distribution function based on current status data etc.

Learning Outcomes

At the end of the day participants should have some new ideas on handling biased sampling problems and survival data. This course is particularly helpful for those who are interested in learning some "theoretical results" and some "applied problems". The accompanied R programs will be discussed.

Background for the Instructor



Jing Qin is a Mathematical Statistician at the Biostatistics Research Branch in the National Institute of Allergy and Infectious Diseases. Dr. Qin's research interests include empirical likelihood method, case-control study, length bias sampling, econometrics, survival analysis, missing data, causal inference, genetic mixture models, generalized linear models, survey sampling and microarray data analysis. He is the author of "Biased sampling, over-identied parametric problems and beyond" (Springer, 2017). He was elected as a Fellow of the American Statistical Association in 2006.



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