

SCIENTIFIC PROGRAM

The Third Erich L. Lehmann Symposium
May 16 - 19, 2007
Rice University

Symposium Chair and Organizer

Javier Rojo
Statistics Department, MS-138
Rice University
6100 Main Street
Houston, TX 77005

Plenary Speakers

Erich L. Lehmann University of California, Berkeley	<i>Some history of optimality</i>
Lawrence D. Brown The Wharton School University of Pennsylvania	<i>A unified view of regression, shrinkage, empirical Bayes, hierarchical Bayes, and random effects</i>
James O. Berger Duke University	<i>Some recent developments in Bayesian model selection</i>
Rodrigo Bañuelos Purdue University	<i>Isoperimetric bounds for Lévy processes</i>
Peter J. Bickel University of California, Berkeley	<i>The collapse of particle filters</i>
Stephen M. Stigler University of Chicago	<i>Karl Pearson and testing statistical hypotheses</i>
Peter J. Huber	<i>On the non-optimality of optimal procedures</i>
Willem R. van Zwet University of Leiden	<i>Statistics and the law: the case of the nonchalant nurse</i>

Invited Session Scientific Committee

Javier Rojo, Chair	Rice University
Jane Ling Wang	University of California, Davis
Rudy Guerra	Rice University
Juliet P. Shaffer	University of California, Berkeley
Wei-Yin Loh	University of Wisconsin, Madison
Peter J. Bickel	University of California, Berkeley
Kjell Doksum	University of Wisconsin, Madison
Yongzhao Shao	New York University
Demissie Alemayehu	Pfizer and Columbia University
James O. Berger	Duke University and SAMSI

Invited Sessions

Young Investigators

Javier Rojo, Organizer

Yolanda Muñoz Maldonado, Chair

Brisa N. Sánchez

University of Michigan

Residual-based diagnostics for structural equation models

Yolanda Muñoz Maldonado

UT Health Sc Center Houston

Penalized least squares and frequentist and Bayesian mixed-effects models

Farinaz Koushanfar

Rice University

How challenging is the data set?

Statistical Problems in the Analysis of Genomic and Magnetic Resonance Imaging Data

Wei-Yin Loh, Chair

Sunduz Keles

University of Wisconsin, Madison

Statistical issues arising in the study of transcription regulation

Shaw-Hwa Lo

Columbia University

A method toward mapping of common traits

Young K. Truong

UNC, Chapel Hill

Spatio-temporal modeling for fMRI data

Optimality in Bioinformatics: Theory vs Practice

Rudy Guerra, Chair

David Dahl

Texas A&M University

Simultaneous inference for multiple testing and clustering via Dirichlet process mixture models

Chad Shaw

Baylor College of Medicine

Using annotations in the analysis of genome scale data

Rudy Guerra

Rice University

Incorporating biological knowledge in gene expression analysis

Regularized Methods of Classification and Estimation of Nonparametric Regression and Covariance Matrices When Data is High Dimensional

Peter J. Bickel, Chair

Wei-Yin Loh

University of Wisconsin

Regression and variable selection in large p , small n problems

Debashis Paul

University of California, Davis

Principal component analysis for structured high-dimensional data

Ya'acov Ritov

Hebrew University

Consistent learning methods are approximately local

Probability, Levy Process, and Applications

Javier Rojo, Organizer
Rodrigo Bañuelos, Chair

Dennis Cox *Multiscale models for chemical reaction processes*
 Rice University

Davar Khoshnevisan *On some applications of stable processes*
 University of Utah

José Enrique Figueroa *Non-parametric estimation for some models driven
 by Levy processes*
 Purdue University

Multiplicity: Developments and Current Issues

Juliet P. Shaffer, Chair

Charles Lewis *Bayesian decision theory for multiple
 comparisons*
 Fordham University

Helmut Finner *Testing for equivalence in k sample models*
 Deutsches Diabetes-Zentrum

Ajit C. Tamhane *A mixture model approach to estimating the
 number of true null hypotheses and adaptive
 control of FDR*
 Northwestern University

Recent Advances in Non- and Semi-parametric Modeling

Jane-Ling Wang, Chair

Kjell Doksum *Semi-parametric models based on transformations
 and extremes*
 University of Wisconsin

Xihong Lin *Statistical challenges in analyzing mass
 spectrometry proteomic data*
 Harvard University

Naisyin Wang *Analysis of hierarchical biomedical data using
 semiparametric models*
 Texas A&M University

***Statistical Inference for Population Substructures via Clustering,
 Mixture Models and Other Approaches***

Demissie Alemayehu, Yongzhao Shao, Organizers
Yongzhao Shao, Chair

Bruce G. Lindsay *Modal inference: halfway between clustering
 and mixture analyses*
 Pennsylvania State University

Yuwu Xu *A limit theory for likelihood ratio test under
 unidentifiability for general dependent processes*
 Fordham University

Yongzhao Shao *Recent developments in likelihood theory with
 applications to testing homogeneity in finite
 mixture models and other models*
 New York University

Semiparametric Models, Longitudinal Survival Data, False Discovery Rate, and Brain fMRI

Kjell Doksum, Chair

Jane-Ling Wang *Semi-parametric analysis of longitudinal data truncated by event-time*
University of California at Davis

Kam-Wah Tsui and Shijie Tang *Simultaneous testing of multiple hypotheses using generalized p-values*
University of Wisconsin, Madison

Chunming Zhang *Semi-parametric detection of significant activation for brain fMRI*
University of Wisconsin, Madison

Multiple Testing and Subgroup Analysis

James O. Berger, Chair

Juliet P. Shaffer *Multiplicity and subgroup analysis*
University of California

Peter Mueller *The optimal discovery procedure and Bayesian decision rules*
U.T. M.D. Anderson Cancer Center

M. J. Bayarri *Bayesian and frequentist handling of multiple U. testing*
Valencia, Duke U. and SAMSI

CONTRIBUTED PAPERS

Nancy L. Glenn, University of South Carolina, Columbia: *The GEM Algorithm*

Richard C. Ott, Mesa State College: *On the Operating Characteristics of Some Non-parametric Methodologies for the Classification of Distributions by Tail Behavior*

Xiaohui Wang, University of Texas-Pan American: *Classifications of Proteomic Mass Spectra and Other Curve Data*

Xiaohu Li, School of Mathematics and Statistics, Lanzhou University, People's Republic of China: *Stochastic Comparison on Conditional Order Statistics - Some New Results*

Robert Mnatsakanov, West Virginia University, *Some Asymptotic Properties of Varying Kernel Density Estimator*

Monnie McGee, Southern Methodist University, *A Distribution Free Summarization Method for Affymetrix GeneChip[®] Arrays*

Changxiang Rui, University of Arkansas, *Point and Block Prediction in Log-Gaussian Random Fields: The Non-constant Mean Case*

Qiang Zhao, Texas State University, San Marcos, *Survival Analysis of Microarray Gene Expression Data Using Correlation Principal Component Regression*

Sahasini Subba Rao, Texas A&M University, *Normalised Least-Squares Estimation in Time-Varying arch Models*

Santanu Chakraborty, UT Pan-American, *Parametric Inference on Zero-Inflated Poisson Distribution and Its Variants*

Hongxiao Zhu, Rice University, *A Functional Generalized Linear Model with Application to Cervical Pre-cancer Diagnosis Using Fluorescence Spectroscopy*

Xiaowei Wu, Rice University, *Some Estimation and Hypothesis Testing Problems in Fluctuation Analysis*

John Fresen, University of Missouri - Columbia, *On the Definition of Weak Convergence of a Sequence of Random Elements*

Victor De Oliveira, UT San Antonio, *Objective Bayesian Analysis of Spatial Data with Measurement Error*

Pang Du, Virginia Tech, *Smoothing Spline Frailty Model*

Graciela Gonzalez, CIMAT, *Some Important Issues in Inference under Certain Types of Singularities*

The Third Erich L. Lehmann Symposium
Javier Rojo, Organizer/Chair
Invited Program – All talks are in McMurtry Auditorium Duncan Hall 1055

	Wednesday May 16	Thursday May 17	Friday May 18	Saturday May 19
7:30	Breakfast 7:30 – 8:45 Duncan Hall	Breakfast Duncan Hall	Breakfast Duncan Hall	Breakfast Duncan Hall
8:45 - 9:00	Opening Remarks Plenary Speaker Erich L. Lehmann	Classification, nonparametric regression: high dimensional data Wei-Yin Loh Debashis Paul Ya'aqov Ritov	Recent Advances in Non and Semiparametric Modeling Kjell Doksum Xihong Lin Naisyin Wang	Multiple Testing and Subgroup analysis Juliet P. Shaffer Peter Mueller Susie Bayarri
10:00	Young Investigators Brisa N. Sánchez	Coffee Break 10:30 – 10:50	Coffee Break	Coffee Break
10:30 – 10:50	Yolanda Muñoz Maldonado Farinaz Koushanfar	Plenary Speaker 10:50 Rodrigo Bañuelos	Plenary Speaker 10:50 Lawrence D Brown	Plenary Speaker 10:50 Peter J Huber
	Lunch 12:00 – 1:15 Duncan Hall	Lunch 11:50 – 1:15 Duncan Hall	Lunch 11:50 – 1:15 Duncan Hall	Plenary Speaker 11:50 Willem van Zwet
1:15 – 3:00	Genomics and magnetic resonance Sunduz Keles Shaw-Hwa Lo Young K. Truong	Probability, Lévy Processes and Applications Dennis D. Cox Davar Khoshnevisan Jose Enrique Figueroa	Inference for Substructures: Clustering, Mixtures, ... Bruce G. Lindsay Yuewu Xu Yongzhao Shao	CLOSING LUNCHEON DUNCAN HALL 1:00 – 2:30
3:00 – 3:20	Coffee Break	Coffee Break	Coffee Break	
3:20 - 4:20	Plenary Speaker Peter J Bickel	Plenary Speaker Stephen M Stigler	Plenary Speaker James O Berger	
4:20 – 6:05	<u>Optimality in Bioinformatics:</u> Theory vs Practice David B. Dahl Chad Shaw Rudy Guerra	<u>Multiplicity: Developments and current issues</u> Charles Lewis Helmut Finner Ajit C. Tamhane	<u>Semiparametrics, longitudinal survival, false discovery, FMRI</u> Jane-Ling Wang Kam-Wah Tsui Chunning Zhang	
	<u>CONTRIBUTED TALKS</u> 6:05 – 7:05	Banquet Ballroom Ley Student Ct Cash Bar/Music 6:45 – 7:30 Dinner/Music 7:30 – 10:00	<u>CONTRIBUTED TALKS</u> 6:05 – 7:05	