

Author(s)	Poster Title
Ackermann, Marit; Sikora-Wohlfeld, Weronika; Beyer, Andreas	Mapping of dynamic expression QTL
Aravena, Andrés; Guziolowski, Carito; Siegel, Anne; Maass, Alejandro	Using Mutual Information and Answer Set Programming to refine Acidithiobacillus ferrooxidans whole genome transcriptional regulation network
Becker, Benjamin	Ein Zweistichprobentest für Split-Plot-Designs mit zufällig fehlenden Werten bei unbekanntem Kovarianzmatrizen
Berres, Manfred; Ruth, Volland; Irene, Beck; Andreas, Monsch	Orthogonal regression – an alternative method to ordinary least squares
Böhning, Dankmar	Mantel-Haenszel Estimation for a General Risk Ratio
Boldt, Sonja; Knops, Katja; Kriehuber, Ralf; Wolkenhauer, Olaf	Identification of gene-based biosignatures for radiation biodosimetry
Enders, Dirk; Schill, Walter	Die Performance von Regression Calibration und SIMEX in einer Ereigniszeitanalyse mit Messfehlern in den Kovariablen unter einem komplexen Fehlermodell
Estaghvirou, Boubacar; Piepho, Hans-Peter; Joseph, O. Ogutu,; Schulz-Streeck, Torben; Technow, Frank	Efficient computation of ridge-regression BLUP in genomic selection in plant breeding
Feilke, Martina C.; Schmid, Volker J.	Bayesian Analysis of FRAP-Images with Mixed-Effect Priors
Fermin, Yessica; Wieczorek, Jakob; Ickstadt, Katja	Network Analysis with Nonparametric Bayesian Networks
Fuhlbrück, Frederike; Rucker, Simon; Geinets, Anna; Senkov, Natalja; Kunde, Steven; Wascher, Lan; Allignol, Arthur; Meister, Reinhard	Display of missing values Surveillance of A(H1N1) vaccination in pregnancy
Fuhlbrück, Frederike; Rucker, Simon; Geinets, Anna; Senkov, Natalja; Wascher, Lan; Kunde, Steven; Allignol, Arthur; Meister, Reinhard	Display of complete datasets Surveillance of A(H1N1) vaccination in pregnancy
Gawrych, Katarzyna; Lotz, Anne; Kendzia, Benjamin; Weiß, Tobias; Brüning, Thomas; Pesch, Beate	Statistische Analysemethoden für linkszensierte Variablen mit Anwendung auf Simulations- und Messdaten
Gondan, Matthias; Scherag, André	A nonparametric test for weak stochastic dominance
Hasan, Hayan	A modified partial least squares approach

Author(s)	Poster Title
Igl, Bernd-Wolfgang; Girbig, Dorothee; Hischke, Sandra; Keller, Karsten; Prella, Katja; Patchev, Vladimir; Vonk, Richardus	A new statistical strategy to analyze circadian body temperature rhythms in a "hot flush" animal model
Ingel, Katharina; Jahn-Eimermacher, Antje	Schätzen der Varianzinflation im Andersen-Gill-Modell
Kabisch, M; Dünnebier, T; Melchior, F; Pesch, B; Brüning, T; Rabstein, S; Ko, Y; Baisch, C; Fischer, H-P; Justenhoven, C; Brauch, H; Harth, V; Hamann, U; Lorenzo Bermejo, J	Refinement of Genetic Associations using Genotype Imputation with Data from 1000 Genomes Project
Kesselmeier, Miriam; Longerich, Thomas; Geffers, Robert; Lorenzo Bermejo, Justo	Comparison of count models regarding their ability to capture the relationship between genomic instability and methylation in human hepatocellular carcinoma
Klotsche, Jens; Liedmann, Ina; Niewerth, Marina; Minden, Kirsten	Binary regression: The measures of total gain in positive and negative predictive value in evaluating the usefulness of a risk prediction model
Koch, Raphael; Gerß, Joachim; Köpcke, Wolfgang; Faldum, Andreas	Improvement of a Bayesian model: Combining results from adult and pediatric clinical trials for planning and analyzing a new pediatric trial.
Köllmann, Claudia; Bornkamp, Björn; Fried, Roland; Ickstadt, Katja	Penalized unimodal spline regression
Kunz, Cornelia Ursula; Friede, Tim; Parsons, Nicholas; Todd, Susan; Stallard, Nigel	Comparison of Methods for Treatment Selection in Seamless Phase II/III Trials Incorporating Information on Short-term Endpoints
Kunz, Cornelia Ursula; Kieser, Meinhard	Schätzung von sekundären Endpunkten in mehrstufigen Phase-II-Studien in der Onkologie
Lorenz, Eva; Jenkner, Carolin; Sauerbrei, Willi; Becher, Heiko	Analytical dose-response functions for univariate and bivariate distribution classes and application of a modified fractional polynomial procedure.
Mayer, Benjamin; Hegele, Lisa	Die Cluster-Delta-Methode zur Ersetzung fehlender Werte in klinischen Verlaufsstudien

Author(s)	Poster Title
Miosga, Pascal; Chung, Tje-Lin; Ackermann, Hanns; Herrmann, Eva; Helm, Werner	A new approach towards simulation of the Rule of Ten for a Cox Regression
Moecks, Joachim	Non-parametric two-sample tests: tuning of raw data before ranking
Möhring, Jens; Piepho, Hans-Peter	Efficiency of augmented p-rep designs
Neumann, Konrad; Pieper, Robert	Finding relevant metabolites by Partial Least Squares (PLS) analysis
Peil, Barbara; Meder, Benjamin; Stoll, Monika; Katus, Hugo A.; Lorenzo Bermejo, Justo	Taking advantage of genotype imputation to detect the association between known risk variants and disease subphenotypes in dilated cardiomyopathy
Rücker, Gerta	All in one: Presenting simulation results in a single plot
Schaffrath Rosario, Angelika; Stolzenberg, Heribert	Daten für alle. Public Use Files des Robert Koch-Instituts und ihre Auswertung.
Scherb, Hagen; Voigt, Kristina	Fehlbildungen, Säuglingssterblichkeit, Schilddrüsenkrebs und das Geschlechtsverhältnis bei der Geburt vor und nach der Tschernobyl-Katastrophe
Schmidtman, Irene	Wie entsteht Wissen – ein neues interdisziplinäres Lehrprojekt an der Johannes Gutenberg-Universität Mainz
Schneider, Michael; Frömke, Cornelia; Zapf, Antonia	High-Performance in R: Bootstrap-Verfahren als robuste Schätzer der Area under the ROC-Curve (AUC) unter der Verwendung von C-kompilierten Prozeduren
Schwender, Holger; Bowers, Katherine; Fallin, M. Danielle; Ruczinski, Ingo	Detecting SNP Interactions in Case-Parent Trio Studies with trioFS
Simang, Michael; Mansmann, Ulrich	Assessing the use of a modified AIC for the selection between GEE models
Stange, Jens; Dickhaus, Thorsten	Über die effektive Anzahl simultan durchzuführender Chi-Quadrat-Tests
Unger, Sigrun	Type II error in dose proportionality studies
Vock, Michael	The horse kick data revisited
Weyer, Veronika; Zwiener, Isabella; Hieke, Stefanie; Butterbach, Katja; Nieters, Alexandra; Schumacher, Martin; Binder, Harald	Developing subgroup SNP signatures by weighted regression for case-control data