

Publication List

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Publications

Journal Papers

1. Chen, Z. and Y. Jiang (2016). An iterated alternate sequential procedure for simultaneous feature selection and precision matrix estimation in sparse high-dimensional multivariate regression models. Submitted.
2. Jiang, Y. and Z. Chen (2016). A Sequential Scaled Pairwise Selection Approach to Edge Detection in Nonparanormal Graphical Models. *Canadian Journal of Statistics* **44(1)**, 25-43.
3. Y. He and Z. Chen (2016). A sequential procedure for feature selection with linear interactive models in ultra-high dimensional feature space. *Annals of the Institute of Statistical Mathematics*, **68(1)**, 155-180.
4. S. Luo, J. Xu and Z. Chen (2015). Extended Bayesian Information Criterion in the Cox Model with a High-dimensional Feature Space. *Annals of the Institute of Statistical Mathematics*, **67** (2), 287-311.
5. S. Luo and Z. Chen (2014). Edge Detection in Sparse Gaussian Graphical Models. *Computational Statistics and Data Analysis* **70** 138-152.
6. S. Luo and Z. Chen (2014). Sequential Lasso cum EBIC for feature selection with ultra-high dimensional feature space. *Journal of the American Statistical Association*, **109**(507), 1229-1240.
7. S. Luo and Z. Chen (2013). Extended BIC for linear regression models with diverging number of parameters and high or ultra-high feature spaces. *Journal of Statistical Planning and Inference*, **143** 494-504.

8. S. Luo and Z. Chen (2013). Selection consistency of EBIC for GLIM with non-canonical links and diverging number of parameters. *Statistics and Its Interface*. **6**, 275-284.
9. J. Chen and Z. Chen (2012). Extended BIC for small- n -large- P sparse GLM. *Statistica Sinica* **22** 555-574.
10. J. Zhao and Z. Chen (2012). A Two-stage penalized logistic regression approach to case-control genome-wide association studies *Journal of Probability and Statistics*, vol. 2012, Article ID 642403, 15 pages, 2012. doi:10.1155/2012/642403.
11. J. Joo, M. Kwak, Z. Chen and G. Zheng (2010). Tutorial in biostatistics: Efficiency robust statistics for genetic linkage and association studies under genetic model uncertainty. *Statistics in Medicine* **29**(1):158-80.
12. Z. Chen and W. Cui (2010). A two-phase procedure for QTL mapping with regression models. *Theoretical and Applied Genetics*. **121**:363-372.
13. Z. Chen, J. Chen. (2009). Tournament Screening cum EBIC for Feature Selection with High Dimensional Feature Spaces. *Science in China, Series A*, **52**(6): 1327-1341.
14. W. Li and Z. Chen (2009). Multiple interval QTL mapping for trait distribution with a spike. *Genetics*, **182**:337-342.
15. W. Li and Z. Chen (2009). A New Approach to Interval Mapping of Quantitative Trait Loci Using Sib-Pairs. *Statistical Methodology*. **5**, 251-261.
16. Z. Chen and J. Liu. (2009) Mixture Generalized Linear Models for Multiple Interval Mapping of Quantitative Trait Loci in Experimental Crosses. *Biometrics*, **65**: 470-477.
17. J. Chen and Z. Chen. (2008). Extended Bayesian Information Criteria for Model Selection with Large Model Space. *Biometrika*, **95**: 759-771.

18. Z. Chen, J. Liu, L. Shen and Y. G. Wang.(2008). General ranked set sampling for efficient treatment comparisons. *Statistica Sinica*, **18**,91-104.
19. Z. Chen. (2007). Ranked Set Sampling: Its Essence and New Applications. *Environmental and Ecological Statistics*. **14**, 355-363.
20. G. Zheng, K. Ghosh, Z. Chen and Z. Li. (2006). Extreme rank selections for linkage analysis of quantitative trait loci using selected sib-pairs. *Annals of Human Genetics*, **70**, 857-866.
21. H. Chen and Z. Chen (2005). Asymptotic properties of the remedian. *Journal of Nonparametric Statistics*. **17**, 155-165.
22. Z. Chen and S. Zheng (2005). Lifetime distribution based degradation analysis. *IEEE: Transactions on Reliability*. **54** 3-10.
23. G. Zheng and Z. Chen. (2005). Comparison of maximum statistics for hypothesis testing when a nuisance parameter presents only under the alternative. *Biometrics*, **61**, 254-258.
24. Z. Chen and G. Zheng. (2005) Exact robust tests for detecting candidate-gene association in case-parents trio design. *Journal of Data Science*, **3** 19-33.
25. Z. Chen (2005). The full EM algorithm for the MLEs of QTL effects and positions and their estimated variances in multiple interval mapping. *Biometrics*, **61**, 474-480.
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28. Z. Chen and Y. Wang.(2004) Efficient regression analysis with ranked set sampling. *Biometrics*, **60**, 997-1004.

29. G. Zheng and Z. Chen. (2004) Robust tests for candidate-gene association using case-parents trios with a multi-allelic marker. *Biometrical Journal*, **46**, 606-611.
30. Y. G. Wang, Z. Chen and J. Liu (2004). General ranked set sampling with cost considerations. *Biometrics* **60**, 556-561.
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32. G. Zheng, Z. Chen and Z. Li (2003). Test for candidate-gene association using case-parents design when the genetic model is unknown. *Annals of Human Genetics* **67**, 589-597.
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34. Z. Chen (2003). Analysis of component reliability of k -out-of- n systems with censoring, *Journal of Statistical Planning and Inference* **116** 305-315.
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56. Y. Zhang and Z. Chen (1982). Optimal supplementary experiments, *Wuhan Daxue Xuebao*. (Chinese, English Summary.)
57. Z. Chen and J. Liu. Multiple Interval Mapping for Ordinal Traits Using Mixture GLIM. *Manuscript*.

Books

1. Z. Chen (2014). *Statistical Methods for QTL Mapping*. ISBN 978-1-4398-6830-0. Chapman & Hall/CRC, New York.
2. Z. Chen, Z. D. Bai and B. K. Sinha (2004). *Ranked Set Sampling: Theory and Applications*. ISBN 0-387-40263-2. Springer-Verlag, New York.
3. Z. Chen, J.T Zhang and F. Hu (2008). *Advances in Statistics*. World Scientific.