

# Short CV

## James G. Booth

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### Contact information

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### Profile

I am currently a Professor in the Department of Biological Statistics and Computational Biology (BSCB) in the College of Agriculture and Life Sciences at Cornell University. I visited the Department of Operations Research and Information Engineering at Cornell in 2003, and was hired in BSCB the following year. From 1987 to 2003 I was a faculty member in the Department of Statistics at the University of Florida. During that period I spent two years as a Research Fellow at the Australian National University, and one year at Colorado State University.

My main research interests involve basic statistical methodology including: the bootstrap and Monte Carlo methods, clustering, exact inference, mixed models, generalized linear models, and applications in bioinformatics. I have taught a variety of courses at Cornell including the second semester of a statistical methods sequence for graduate students from a wide variety of disciplines, as well as core courses for statistics undergraduates, professional masters students, and Ph.D. students in the Fields of Statistics. As a faculty member in BSCB I am also actively involved in the Cornell Statistical Consulting Unit which provides advice on statistical issues to faculty and students at Cornell.

### Education

PhD	Statistics	1987	University of Kentucky
MS	Statistics	1984	University of Kentucky
MSc	Statistics	1982	University of Leeds, England
BSc	Mathematics	1981	University of Leeds, England

### Positions Held

Professor	2004 –present	Cornell University
Visiting Professor	2003 – 2004	Cornell University
Professor	2000 – 2004	University of Florida
Visiting Professor	1996 – 1997	Colorado State University
Associate Professor	1993 – 2000	University of Florida
Research Fellow	1990 – 1992	Australian National University
Assistant Professor	1987 – 1993	University of Florida
Instructor	1986 – 1987	Transylvania University, KY

### Departmental Service

- *Director of Graduate Studies*, Field of Statistics, Cornell University, 2017–present.
- *Chair*, Department of Biological Statistics and Computational Biology, College of Agriculture and Life Sciences, Cornell University, 2007–2016

- *Director of Undergraduate Studies*, Biometry and Statistics, College of Agriculture and Life Sciences, Cornell University, 2011–2016
- *Acting Chair*, Department of Statistics, University of Florida, Fall 2002
- *Associate Chair*, Department of Statistics, University of Florida, 2001–2003
- *Graduate Coordinator*, Department of Statistics, University of Florida, 1995–1996, 1997–2001

## Professional Service

- *External Review Committee*: Department of Statistics, Southern Methodist University, Nov. 2012.
- *Executive Committee*: Statistical Modelling Society, 2006–2012.
- *Associate Editor*: Journal of Statistical Planning and Inference, 2012–2014, Journal of the American Statistical Association, 2006–2012, Statistical Modelling: 2001–, ANZ Journal of Statistics: 2001–2008, Journal of the Royal Statistical Society, Series B: 2001–2005, Computational Statistics: 1996–2002
- *Book Review Board*: Journal of the American Statistical Association: 1999–2002
- *NSF Review Panel*: Program in Statistics: Dec. 10–12, 2001 and Feb. 6–8, 2013.

## Graduate Students Supervised

- Kelson Zawack, PhD in Computational Biology, Cornell University, May 2017 (Co-adviser with Yrjo Grohn). Postdoctoral Associate, Department of Biostatistics, Yale University.
- Irina Gaynanova, PhD, Cornell University, May 2015 (Co-adviser with Martin Wells). Assistant Professor, Department of Statistics, Texas A&M University.
- Muting Wan, PhD, Cornell University, January 2015. Data Scientist, New York Life.
- Caitlin Cunningham, Cornell University, August 2012. Assistant Professor, LeMoyne College, Syracuse.
- Haim Bar, PhD, Cornell University, January 2012. Assistant Professor, Department of Statistics, University of Connecticut.
- Kirsten Eilertson, PhD, Cornell University, August 2011. Research Associate, Department of Statistics, Pennsylvania State University.
- Matthias Kormacksson, PhD, Cornell University, August 2009. Research Scientist, IBM Research, Rio de Janeiro.
- Vadim Zipunikov, PhD, Cornell University, January 2008. Assistant Professor, Department of Biostatistics, Johns-Hopkins University.
- David Hitchcock, PhD, August 2004. Associate Professor at University of South Carolina. (Co-advisor George Casella)
- Bernhard Klingenberg, PhD, August 2004. Associate Professor, Williams College. (Co-advisor Alan Agresti)
- Brian Caffo, PhD, August 2001. Professor, Department of Biostatistics, Johns-Hopkins University.

- Wolfgang Jank, PhD, August 2001. Anderson Professor of Global Management, Department of Information Systems and Decision Sciences, University of South Florida.
- Glen Hartless, PhD, August 2000. CIA Analyst in Washington D.C. (Co-advisor: Ramon Littell)
- Somnath Sarkar, PhD, August 1996. Statistician at Eli-Lilly, Indianapolis. (Co-advisor: M. Ghosh)

## Grants

- co-PI (PI Haim Bar, co-PI Martin Wells), NSF DMS-1611893, “Variable selection when  $p \gg N$  - beyond the linear regression and normal errors model” 2016-2019.
- PI (co-PI Martin Wells), NSF DMS-1208488, “Models and computational strategies in statistical bioinformatics” 2012-2014.
- PI, NSA Grant 08C-016, “24th International Workshop on Statistical Modeling” 2008-2009.
- PI, NSF DMS-0805865, “Applications and computational issues involving generalized linear and mixed models”, 2008-2010.
- Co-PI, NSF Grant DMS-0405543, “Cluster analysis, predictive distributions, and stochastic search algorithms,” 2004-2008. PI George Casella, Department of Statistics, University of Florida.
- Co-PI, NSF Grant DMS-0072827, “Combining EM and Monte Carlo to maximize intractable likelihood functions,” 2000-2004.
- PI, NSF Grant DMS-9813374, “NSF/CBMS Regional Conference in the Mathematical Sciences - Generalized Linear Mixed Models and Related Topics - June 8-12, 1999.”
- PI, NSF Grant DMS-93010836, “Some new bootstrap methods for sample surveys,” 1993-1996.

## Selected Publications

- [1] BAR, H. Y., BOOTH, J. G. & WELLS, M. T. (2014). A bivariate model for simultaneous testing in bioinformatics data. *Journal of the American Statistical Association* **109**, 537–547.
- [2] BOOTH, J. G. & BUTLER, R. W. (1999). An importance sampling algorithm for exact conditional tests in log-linear models. *Biometrika* **86**, 321–332.
- [3] BOOTH, J. G., BUTLER, R. W. & HALL, P. (1994). Bootstrap methods for finite populations. *Journal of the American Statistical Association* **89**, 1282–1289.
- [4] BOOTH, J. G., CAPANU, M. & HEIGENHAUSER, L. (2005). Exact conditional p-value calculation for the quasi-symmetry model. *Journal of Computational and Graphical Statistics* **14**, 716–725.
- [5] BOOTH, J. G., CASELLA, G. & HOBERT, J. P. (2008). Clustering using objective functions and stochastic search. *Journal of the Royal Statistical Society B* **70**, 119–139.
- [6] BOOTH, J. G., FEDERER, W. T., WELLS, M. T. & WOLFINGER, R. (2009). A multivariate variance component model for analysis of covariance in designed experiments. *Statistical Science* **24**, 223–237.

- [7] BOOTH, J. G. & HALL, P. (1993). Bootstrap confidence regions for functional relationships in errors-in-variables models. *Annals of Statistics* **21**, 1780–1791.
- [8] BOOTH, J. G. & HALL, P. (1994). Monte Carlo approximation and the iterated bootstrap. *Biometrika* **81**, 331–340.
- [9] BOOTH, J. G., HALL, P. & WOOD, A. T. A. (1993). Balanced importance resampling for the bootstrap. *Annals of Statistics* **21**, 286–298.
- [10] BOOTH, J. G. & HOBERT, J. P. (1998). Standard errors of prediction in generalized linear mixed models. *Journal of the American Statistical Association* **93**, 262–272.
- [11] BOOTH, J. G. & HOBERT, J. P. (1999). Maximizing generalized linear mixed model likelihoods with an automated Monte Carlo EM algorithm. *Journal of the Royal Statistical Society B* **61**, 265–285.
- [12] BOOTH, J. G. & PRESNELL, B. (1998). Allocation of Monte Carlo resources for the iterated bootstrap. *Journal of Computational and Graphical Statistics* **7**, 92–112.
- [13] CAFFO, B. S. & BOOTH, J. G. (2001). A Markov chain Monte Carlo algorithm for approximating exact conditional tests. *Journal of Computational and Graphical Statistics* **10**, 730–745.
- [14] CAFFO, B. S. & BOOTH, J. G. (2003). Monte Carlo conditional tests for log-linear and logistic models: a survey of current methodology. *Statistical Methods in Medical Research* **12**, 1–15.
- [15] CAFFO, B. S., BOOTH, J. G. & DAVISON, A. C. (2002). Empirical sup rejection sampling. *Biometrika* **89**, 745–754.
- [16] GAYNANOVA, I., BOOTH, J. G. & WELLS, M. T. (2016). Penalized versus constrained generalized eigenvalue problems. *Journal of Computational and Graphical Statistics* Posted online 06 Apr 2016.
- [17] GAYNANOVA, I., BOOTH, J. G. & WELLS, M. T. (2016). Simultaneous sparse estimation of canonical vectors in the  $p \gg n$  setting. *Journal of the American Statistical Association* **111**, 696–706. Published online 16 Apr 2015.
- [18] HARTLESS, G. L., BOOTH, J. G. & LITTELL, R. C. (2003). Local influence of predictors in multiple linear regression. *Technometrics* **45**, 326–332.
- [19] HITCHCOCK, D. B., CASELLA, G. & BOOTH, J. G. (2006). Improved estimation of dissimilarities by smoothing functional data. *Journal of the American Statistical Association* **101**, 211–222.
- [20] JANK, W. & BOOTH, J. G. (2003). Efficiency of Monte Carlo EM and simulated maximum likelihood in generalized linear mixed models. *Journal of Computational and Graphical Statistics* **12**, 214–229.
- [21] KORMAKSSON, M., BOOTH, J. G., FIGUEROA, M. E. & MELNICK, A. (2012). Integrative model-based clustering of microarray methylation and expression data. *Annals of Applied Statistics* **6**, 1327–1347.

- [22] ZIPUNNIKOV, V., BOOTH, J. G. & YOSHIDA, R. (2009). Table counting using the saddlepoint approximation. *Journal of Computational and Graphical Statistics* **18**, 915–929.