

Daniel A Henderson

List of Publications by Year in descending order

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14
papers

350
citations

1040056

9
h-index

1199594

12
g-index

14
all docs

14
docs citations

14
times ranked

379
citing authors

#	ARTICLE	IF	CITATIONS
1	On Bayesian inference for the Extended Plackett-Luce model. <i>Bayesian Analysis</i> , 2021, -1, .	3.0	0
2	Revealing Subgroup Structure in Ranked Data Using a Bayesian WAND. <i>Journal of the American Statistical Association</i> , 2020, 115, 1888-1901.	3.1	3
3	Emulation of Utility Functions Over a Set of Permutations: Sequencing Reliability Growth Tasks. <i>Technometrics</i> , 2018, 60, 273-285.	1.9	3
4	A Comparison of Truncated and Time-Weighted Plackett-Luce Models for Probabilistic Forecasting of Formula One Results. <i>Bayesian Analysis</i> , 2018, 13, .	3.0	13
5	Adaptive, Delayed-Acceptance MCMC for Targets With Expensive Likelihoods. <i>Journal of Computational and Graphical Statistics</i> , 2017, 26, 434-444.	1.7	25
6	Delayed acceptance particle MCMC for exact inference in stochastic kinetic models. <i>Statistics and Computing</i> , 2015, 25, 1039-1055.	1.5	50
7	Regional variations in the European Neolithic dispersal: the role of the coastlines. <i>Antiquity</i> , 2014, 88, 1291-1302.	1.0	9
8	Life Sciences at a Crossroads: Respiratory Transmissible H5N1. <i>Science</i> , 2012, 335, 801-802.	12.6	25
9	Bayesian Calibration of a Stochastic Kinetic Computer Model Using Multiple Data Sources. <i>Biometrics</i> , 2010, 66, 249-256.	1.4	11
10	Bayesian Emulation and Calibration of a Stochastic Computer Model of Mitochondrial DNA Deletions in Substantia Nigra Neurons. <i>Journal of the American Statistical Association</i> , 2009, 104, 76-87.	3.1	60
11	Discussion of "A Bayesian Approach to DNASequencing Segmentation ". <i>Biometrics</i> , 2005, 61, 637-639.	1.4	0
12	A Bayesian Approach to DNA Sequence Segmentation. <i>Biometrics</i> , 2004, 60, 573-581.	1.4	71
13	A compositional segmentation of the human mitochondrial genome is related to heterogeneities in the guanine mutation rate. <i>Nucleic Acids Research</i> , 2003, 31, 6043-6052.	14.5	19
14	Detecting homogeneous segments in DNA sequences by using hidden Markov models. <i>Journal of the Royal Statistical Society Series C: Applied Statistics</i> , 2000, 49, 269-285.	1.0	61