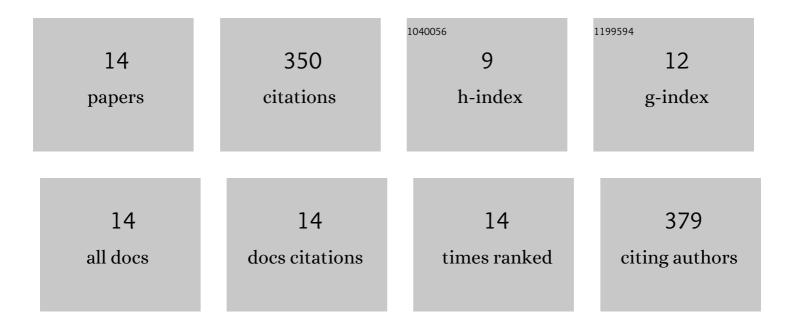
Daniel A Henderson

List of Publications by Year in descending order

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