## Jens Ledet Jensen

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Normalization of Real-Time Quantitative Reverse Transcription-PCR Data: A Model-Based Variance Estimation Approach to Identify Genes Suited for Normalization, Applied to Bladder and Colon Cancer Data Sets. Cancer Research, 2004, 64, 5245-5250.	0.9	5,993
2	Identifying distinct classes of bladder carcinoma using microarrays. Nature Genetics, 2003, 33, 90-96.	21.4	452
3	Bayesian coestimation of phylogeny and sequence alignment. BMC Bioinformatics, 2005, 6, 83.	2.6	169
4	Spatial mixture modeling of fMRI data. Human Brain Mapping, 2000, 11, 233-248.	3.6	117
5	A Dependent-Rates Model and an MCMC-Based Methodology for the Maximum-Likelihood Analysis of Sequences with Overlapping Reading Frames. Molecular Biology and Evolution, 2001, 18, 763-776.	8.9	86
6	Asymptotic normality of the maximum likelihood estimator in state space models. Annals of Statistics, 1999, 27, .	2.6	76
7	Recursions for statistical multiple alignment. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 14960-14965.	7.1	72
8	Synaptic contact number and size in stratum radiatum CA1 of APP/PS1ΔE9 transgenic mice. Neurobiology of Aging, 2009, 30, 1756-1776.	3.1	67
9	Estimation of some aeolian saltation transport parameters: a reâ€analysis of Williams' data. Sedimentology, 1986, 33, 547-558.	3.1	65
10	Probabilistic models of DNA sequence evolution with context dependent rates of substitution. Advances in Applied Probability, 2000, 32, 499-517.	0.7	58
11	On the Laplace Transform of the Lognormal Distribution. Methodology and Computing in Applied Probability, 2016, 18, 441-458.	1.2	55
12	Zinc transporter gene expression is regulated by pro-inflammatory cytokines: a potential role for zinc transporters in beta-cell apoptosis?. BMC Endocrine Disorders, 2009, 9, 7.	2.2	48
13	Statistical Inference in Evolutionary Models of DNA Sequences via the EM Algorithm. Statistical Applications in Genetics and Molecular Biology, 2005, 4, Article18.	0.6	34
14	Markovian approximation to the finite loci coalescent with recombination along multiple sequences. Theoretical Population Biology, 2014, 98, 48-58.	1.1	33
15	Exponential Family Techniques for the Lognormal Left Tail. Scandinavian Journal of Statistics, 2016, 43, 774-787.	1.4	25
16	Probabilistic models of DNA sequence evolution with context dependent rates of substitution. Advances in Applied Probability, 2000, 32, 499-517.	0.7	24
17	Is the â€~improved likelihood ratio statistic' really improved in the discrete case?. Biometrika, 1989, 76, 655-661.	2.4	23
18	Large Deviation and Other Results for Minimum Contrast Estimators. Annals of the Institute of Statistical Mathematics, 1998, 50, 673-695.	0.8	19

JENS LEDET JENSEN

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19	Saddlepoint expansions for sums of Markov dependent variables on a continuous state space. Probability Theory and Related Fields, 1991, 89, 181-199.	1.8	18
20	Validation of the use of DNA pools and primer extension in association studies of sporadic colorectal cancer for selection of candidate SNPs. Human Mutation, 2006, 27, 187-194.	2.5	17
21	Is the `Improved Likelihood Ratio Statistic' Really Improved in the Discrete Case?. Biometrika, 1989, 76, 655.	2.4	15
22	A Large Deviation-Type Approximation for the "Box Class―of Likelihood Ratio Criteria. Journal of the American Statistical Association, 1991, 86, 437-440.	3.1	15
23	Summary Statistics for Endpoint-Conditioned Continuous-Time Markov Chains. Journal of Applied Probability, 2011, 48, 911-924.	0.7	11
24	Applications of Hidden Markov Models for Characterization of Homologous DNA Sequences with a Common Gene. Journal of Computational Biology, 2005, 12, 186-203.	1.6	9
25	The Dynamic Model of Choice for Public Policy Reconsidered: A Formal Analysis With an Application to US Budget Data. Journal of Public Administration Research and Theory, 2016, 26, 226-238.	3.3	9
26	A Large Deviation-Type Approximation for the "Box Class" of Likelihood Ratio Criteria. Journal of the American Statistical Association, 1991, 86, 437.	3.1	8
27	High dimensional classifiers in the imbalanced case. Computational Statistics and Data Analysis, 2016, 98, 46-59.	1.2	8
28	A Simple Derivation of r * for Curved Exponential Families. Scandinavian Journal of Statistics, 1997, 24, 33-46.	1.4	7
29	Bayesian Phylogenetic Inference under a Statistical Insertion-Deletion Model. Lecture Notes in Computer Science, 2003, , 228-244.	1.3	6
30	Higher-Order Asymptotics and Its Application to Testing the Equality of the Examinee Ability Over Two Sets of Items. Psychometrika, 2019, 84, 484-510.	2.1	6
31	A Comparative Distributional Method for Public Administration Illustrated Using Public Budget Data. Journal of Public Administration Research and Theory, 2019, 29, 460-473.	3.3	6
32	Ornstein–Uhlenbeck type processes with non-normal distribution. Journal of Applied Probability, 1999, 36, 389-402.	0.7	6
33	A Simple Derivation of r* for Curved Exponential Families. Scandinavian Journal of Statistics, 1997, 24, 33-46.	1.4	5
34	Markov jump processes with a singularity. Advances in Applied Probability, 2000, 32, 779-799.	0.7	5
35	Summary Statistics for Endpoint-Conditioned Continuous-Time Markov Chains. Journal of Applied Probability, 2011, 48, 911-924.	0.7	4
36	Asymptotic expansions at work. Scandinavian Actuarial Journal, 1995, 1995, 143-152.	1.7	3

Jens Ledet Jensen

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37	A Unifying Framework and Comparison of Algorithms for Nonâ€negative Matrix Factorisation. International Statistical Review, 2020, 88, 29-53.	1.9	3
38	On some problems in the article Efficient Likelihood Estimation in State Space Models by Cheng-Der Fuh [Ann. Statist. 34 (2006) 2026–2068]. Annals of Statistics, 2010, 38, .	2.6	2
39	On a saddlepoint approximation to the Markov binomial distribution. Brazilian Journal of Probability and Statistics, 2013, 27, .	0.4	2
40	Significance evaluation in factor graphs. BMC Bioinformatics, 2017, 18, 199.	2.6	2
41	Asymptotic normality of M-estimators in nonhomogeneous hidden Markov models. Journal of Applied Probability, 2011, 48, 295-306.	0.7	2
42	A Note on the Linear Memory Baum-Welch Algorithm. Journal of Computational Biology, 2009, 16, 1209-1210.	1.6	1
43	Classification Error of the Thresholded Independence Rule. Scandinavian Journal of Statistics, 2015, 42, 32-42.	1.4	1
44	On the Use of Saddlepoint Approximations in High Dimensional Inference. Sankhya A, 2021, 83, 379-392.	0.8	1
45	On the three-dimensional negative binomial distribution. Communications in Statistics - Theory and Methods, 2018, 47, 2314-2326.	1.0	0