

Peter MÃ¼ller

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

553
citations

1478505

6
h-index

1281871

11
g-index

19
all docs

19
docs citations

19
times ranked

500
citing authors

#	ARTICLE	IF	CITATIONS
1	An ANOVA Model for Dependent Random Measures. Journal of the American Statistical Association, 2004, 99, 205-215.	3.1	228
2	Determining the Effective Sample Size of a Parametric Prior. Biometrics, 2008, 64, 595-602.	1.4	188
3	Proteome-wide identification of HSP70/HSC70 chaperone clients in human cells. PLoS Biology, 2020, 18, e3000606.	5.6	43
4	Nonlinear State-Space Models With State-Dependent Variances. Journal of the American Statistical Association, 2003, 98, 377-386.	3.1	34
5	Prior Effective Sample Size in Conditionally Independent Hierarchical Models. Bayesian Analysis, 2012, 7, .	3.0	30
6	Search Algorithms and Loss Functions for Bayesian Clustering. Journal of Computational and Graphical Statistics, 2022, 31, 1189-1201.	1.7	15
7	Editorial: Roles of Hypothesis Testing, p-Values and Decision Making in Biopharmaceutical Research. Statistics in Biopharmaceutical Research, 2021, 13, 1-5.	0.8	6
8	Variance in estimated pairwise genetic distance under high versus low coverage sequencing: The contribution of linkage disequilibrium. Theoretical Population Biology, 2017, 117, 51-63.	1.1	4
9	A Bayesian Random Partition Model For Sequential Refinement and Coagulation. Biometrics, 2019, 75, 988-999.	1.4	3
10	A semi-mechanistic dose-finding design in oncology using pharmacokinetic/pharmacodynamic modeling. Pharmaceutical Statistics, 2022, 21, 1149-1166.	1.3	1
11	Simulation-based sequential design. Pharmaceutical Statistics, 2022, 21, 729-739.	1.3	1
12	Discussion on "Predictively consistent prior effective sample sizes," by Beat Neuenschwander, Sebastian Weber, Heinz Schmidli, and Anthony O'Hagan. Biometrics, 2020, 76, 599-601.	1.4	0