Deepayan Sarkar

List of Publications by Year in descending order

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759233 940533 3,041 18 12 16 citations h-index g-index papers 19 19 19 5822 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	From random matrices to long range dependence. Random Matrices: Theory and Application, 2016, 05, 1650008.	1.1	10
2	A Note on the Construction of Orthogonal Latin Hypercube Designs. Journal of Combinatorial Designs, 2016, 24, 105-111.	0.6	6
3	Single-molecule analysis reveals widespread structural variation in multiple myeloma. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 7689-7694.	7.1	43
4	R Graphics. Handbook of Statistics, 2014, 32, 49-91.	0.6	0
5	Discovery of structural alterations in solid tumor oligodendroglioma by single molecule analysis. BMC Genomics, 2013, 14, 505.	2.8	30
6	An integrative genomic approach identifies p73 and p63 as activators of miR-200 microRNA family transcription. Nucleic Acids Research, 2012, 40, 499-510.	14.5	67
7	Statistical Significance of Optical Map Alignments. Journal of Computational Biology, 2012, 19, 478-492.	1.6	17
8	High-resolution human genome structure by single-molecule analysis. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 10848-10853.	7.1	161
9	Genome-wide MyoD Binding in Skeletal Muscle Cells: A Potential for Broad Cellular Reprogramming. Developmental Cell, 2010, 18, 662-674.	7.0	434
10	iFlow: A Graphical User Interface for Flow Cytometry Tools in Bioconductor. Advances in Bioinformatics, 2009, 2009, 1-3.	5 . 7	7
11	Quality Assessment and Data Analysis for microRNA Expression Arrays. Nucleic Acids Research, 2009, 37, e17-e17.	14.5	45
12	flowCore: a Bioconductor package for high throughput flow cytometry. BMC Bioinformatics, 2009, 10, 106.	2.6	484
13	Extending lattice: using generics and methods to implement new visualization methods within the Trellis framework. Computational Statistics, 2008, 23, 565-572.	1.5	1
14	Lattice., 2008,,.		1,068
15	Coverage and error models of protein-protein interaction data by directed graph analysis. Genome Biology, 2007, 8, R186.	9.6	37
16	Detecting differential gene expression with a semiparametric hierarchical mixture method. Biostatistics, 2004, 5, 155-176.	1.5	526
17	Impairment of the Transcriptional Responses to Oxidative Stress in the Heart of Aged C57BL/6 Mice. Annals of the New York Academy of Sciences, 2004, 1019, 85-95.	3.8	24
18	Age-related impairment of the transcriptional responses to oxidative stress in the mouse heart. Physiological Genomics, 2003, 13, 119-127.	2.3	81