Nigel J Burroughs

List of Publications by Year in descending order

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516710 434195 1,115 33 16 31 citations g-index h-index papers 45 45 45 1575 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Kinetochore tracking in 3D from lattice light-sheet imaging data with KiT. Bioinformatics, 2022, 38, 3315-3317.	4.1	1
2	Kinetochore life histories reveal an Aurora-B-dependent error correction mechanism in anaphase. Developmental Cell, 2021, 56, 3082-3099.e5.	7.0	24
3	Subcellular Euclidean distance measurements with multicolor fluorescence localization imaging in cultured cells. STAR Protocols, 2021, 2, 100774.	1.2	2
4	The Effect of a Linear Tuning between the Antigenic Stimulations of CD4+ T Cells and CD4+ Tregs. Mathematics, 2020, 8, 293.	2.2	6
5	Ensemble-Level Organization of Human Kinetochores and Evidence for Distinct Tension and Attachment Sensors. Cell Reports, 2020, 31, 107535.	6.4	40
6	Bayesian inference of origin firing time distributions, origin interference and licencing probabilities from Next Generation Sequencing data. Nucleic Acids Research, 2019, 47, 2229-2243.	14.5	4
7	Actin turnover ensures uniform tension distribution during cytokinetic actomyosin ring contraction. Molecular Biology of the Cell, 2019, 30, 933-941.	2.1	14
8	Computational modelling predicts substantial carbon assimilation gains for C3 plants with a single-celled C4 biochemical pump. PLoS Computational Biology, 2019, 15, e1007373.	3.2	6
9	Kinesin expands and stabilizes the GDP-microtubule lattice. Nature Nanotechnology, 2018, 13, 386-391.	31.5	81
10	A Hidden Markov Model for Detecting Confinement in Single-Particle Tracking Trajectories. Biophysical Journal, 2018, 115, 1741-1754.	0.5	23
11	Mitotic live-cell imaging at different timescales. Methods in Cell Biology, 2018, 145, 1-27.	1.1	8
12	Modelling the suppression of autoimmunity after pathogen infection. Mathematical Methods in the Applied Sciences, 2018, 41, 8565-8570.	2.3	6
13	Changes in Gene Expression in Space and Time Orchestrate Environmentally Mediated Shaping of Root Architecture. Plant Cell, 2017, 29, 2393-2412.	6.6	49
14	Size matters for single-cell C ₄ photosynthesis in <i>Bienertia</i> . Journal of Experimental Botany, 2017, 68, 255-267.	4.8	12
15	Actomyosin Ring Formation and Tension Generation in Eukaryotic Cytokinesis. Current Biology, 2016, 26, R719-R737.	3.9	95
16	KiT: a MATLAB package for kinetochore tracking. Bioinformatics, 2016, 32, 1917-1919.	4.1	11
17	Regulation of Gene Expression in Shewanella oneidensis MR-1 during Electron Acceptor Limitation and Bacterial Nanowire Formation. Applied and Environmental Microbiology, 2016, 82, 5428-5443.	3.1	59
18	Human kinetochores are swivel joints that mediate microtubule attachments. ELife, $2016, 5, .$	6.0	30

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19	Probing microtubule polymerisation state at single kinetochores during metaphase chromosome motion. Journal of Cell Science, 2015, 128, 1991-2001.	2.0	23
20	Inferring the Forces Controlling Metaphase Kinetochore Oscillations by Reverse Engineering System Dynamics. PLoS Computational Biology, 2015, 11, e1004607.	3.2	29
21	Detection of Diffusion Heterogeneity in Single Particle Tracking Trajectories Using a Hidden Markov Model with Measurement Noise Propagation. PLoS ONE, 2015, 10, e0140759.	2.5	38
22	Super-resolution kinetochore tracking reveals the mechanisms of human sister kinetochore directional switching. ELife, $2015, 4, .$	6.0	20
23	A Virulent Strain of Deformed Wing Virus (DWV) of Honeybees (Apis mellifera) Prevails after Varroa destructor-Mediated, or In Vitro, Transmission. PLoS Pathogens, 2014, 10, e1004230.	4.7	294
24	MosaicSolver: a tool for determining recombinants of viral genomes from pileup data. Nucleic Acids Research, 2014, 42, e123-e123.	14.5	6
25	Correcting for link loss in causal network inference caused by regulator interference. Bioinformatics, 2014, 30, 2779-2786.	4.1	2
26	Solar powered biohydrogen production requires specific localization of the hydrogenase. Energy and Environmental Science, 2014, 7, 3791-3800.	30.8	12
27	Error correction and diversity analysis of population mixtures determined by NGS. PeerJ, 2014, 2, e645.	2.0	4
28	Boltzmann Energy-based Image Analysis Demonstrates that Extracellular Domain Size Differences Explain Protein Segregation at Immune Synapses. PLoS Computational Biology, 2011, 7, e1002076.	3.2	24
29	Matched Sizes of Activating and Inhibitory Receptor/Ligand Pairs Are Required for Optimal Signal Integration by Human Natural Killer Cells. PLoS ONE, 2010, 5, e15374.	2.5	45
30	Close contact fluctuations: The seeding of signalling domains in the immunological synapse. Europhysics Letters, 2007, 77, 48003.	2.0	15
31	Stochasticity and spatial heterogeneity in Tâ€cell activation. Immunological Reviews, 2007, 216, 69-80.	6.0	25
32	Ligand Detection and Discrimination by Spatial Relocalization: A Kinase-Phosphatase Segregation Model of TCR Activation. Biophysical Journal, 2006, 91, 1619-1629.	0.5	51
33	Discriminating self from nonself with short peptides from large proteomes. Immunogenetics, 2004, 56, 311-320.	2.4	39