

Nikolaj Zimic

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

85
citations

1684188

5
h-index

1474206

9
g-index

16
all docs

16
docs citations

16
times ranked

121
citing authors

#	ARTICLE	IF	CITATIONS
1	Computational modelling of genome-scale metabolic networks and its application to CHO cell cultures. <i>Computers in Biology and Medicine</i> , 2017, 88, 150-160.	7.0	24
2	Computational analysis of viable parameter regions in models of synthetic biological systems. <i>Journal of Biological Engineering</i> , 2019, 13, 75.	4.7	12
3	The Ternary Quantum-dot Cellular Automata Memorizing Cell. , 2009, , .		9
4	An adaptive genetic algorithm for parameter estimation of biological oscillator models to achieve target quantitative system response. <i>Natural Computing</i> , 2014, 13, 119-127.	3.0	7
5	Semi-quantitative Modelling of Gene Regulatory Processes with Unknown Parameter Values Using Fuzzy Logic and Petri Nets. <i>Fundamenta Informaticae</i> , 2018, 160, 81-100.	0.4	7
6	Notes on fuzzy cellular automata. <i>Journal of the Chinese Institute of Industrial Engineers</i> , 2000, 17, 469-476.	0.5	6
7	Distributed biological computation: from oscillators, logic gates and switches to a multicellular processor and neural computing applications. <i>Neural Computing and Applications</i> , 2021, 33, 8923-8938.	5.6	6
8	Integration of omics data to generate and analyse COVID-19 specific genome-scale metabolic models. <i>Computers in Biology and Medicine</i> , 2022, 145, 105428.	7.0	5
9	Quantum-dot Field Programmable Gate Array: enhanced routing. , 2006, , .		2
10	Grohar: Automated Visualization of Genome-Scale Metabolic Models and Their Pathways. <i>Journal of Computational Biology</i> , 2018, 25, 505-508.	1.6	2
11	Field-programmable biological circuits and configurable (bio)logic blocks for distributed biological computing. <i>Computers in Biology and Medicine</i> , 2021, 128, 104109.	7.0	2
12	Computational Framework for Modeling Multiple Noncooperative Transcription Factor Binding and Its Application to the Analysis of Nuclear Factor Kappa B Oscillatory Response. <i>Journal of Computational Biology</i> , 2016, 23, 923-933.	1.6	1
13	Initial state perturbations as a validation method for data-driven fuzzy models of cellular networks. <i>BMC Bioinformatics</i> , 2018, 19, 333.	2.6	1
14	Towards Multistate Nanocomputing: The Implementation of a Primitive Fuzzy Controller. , 2008, , .		0
15	Space complexity optimization for nano electronic devices based on evolutionary computation. , 2008, , .		0