

Alberto Marin-Sanguino

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1786573/publications.pdf>

Version: 2024-02-01

19
papers

538
citations

840776

11
h-index

839539

18
g-index

23
all docs

23
docs citations

23
times ranked

635
citing authors

#	ARTICLE	IF	CITATIONS
1	Adaptation to Varying Salinity in <i>Halomonas elongata</i> : Much More Than Ectoine Accumulation. <i>Frontiers in Microbiology</i> , 2022, 13, 846677.	3.5	8
2	Anaplerotic Pathways in <i>Halomonas elongata</i> : The Role of the Sodium Gradient. <i>Frontiers in Microbiology</i> , 2020, 11, 561800.	3.5	6
3	Understanding biochemical design principles with ensembles of canonical non-linear models. <i>PLoS ONE</i> , 2020, 15, e0230599.	2.5	4
4	Editorial: Foundations of Theoretical Approaches in Systems Biology. <i>Frontiers in Genetics</i> , 2018, 9, 290.	2.3	0
5	Revision and reannotation of the <i>Halomonas elongata</i> DSM 2581 T genome. <i>MicrobiologyOpen</i> , 2017, 6, e00465.	3.0	13
6	Osmoregulation in the Halophilic Bacterium <i>Halomonas elongata</i> : A Case Study for Integrative Systems Biology. <i>PLoS ONE</i> , 2017, 12, e0168818.	2.5	49
7	Time Hierarchies and Model Reduction in Canonical Non-linear Models. <i>Frontiers in Genetics</i> , 2016, 7, 166.	2.3	4
8	Design Principles as a Guide for Constraint Based and Dynamic Modeling: Towards an Integrative Workflow. <i>Metabolites</i> , 2015, 5, 601-635.	2.9	7
9	Chemical reaction network approaches to Biochemical Systems Theory. <i>Mathematical Biosciences</i> , 2015, 269, 135-152.	1.9	18
10	Biochemical Pathway Modeling Tools for Drug Target Detection in Cancer and Other Complex Diseases. <i>Methods in Enzymology</i> , 2011, 487, 319-369.	1.0	20
11	A blueprint of ectoine metabolism from the genome of the industrial producer <i>Halomonas elongata</i> DSM 2581 T. <i>Environmental Microbiology</i> , 2011, 13, 1973-1994.	3.8	224
12	Steady-state global optimization of metabolic non-linear dynamic models through recasting into power-law canonical models. <i>BMC Systems Biology</i> , 2011, 5, 137.	3.0	21
13	Flux duality in nonlinear GMA systems: Implications for metabolic engineering. <i>Journal of Biotechnology</i> , 2010, 149, 166-172.	3.8	8
14	Optimization of biochemical systems through mathematical programming: Methods and applications. <i>Computers and Operations Research</i> , 2010, 37, 1427-1438.	4.0	41
15	Metabolic Engineering with power-law and linear-logarithmic systems. <i>Mathematical Biosciences</i> , 2009, 218, 50-58.	1.9	3
16	Optimization of biotechnological systems through geometric programming. <i>Theoretical Biology and Medical Modelling</i> , 2007, 4, 38.	2.1	32
17	Optimization of biochemical systems by linear programming and general mass action model representations. <i>Mathematical Biosciences</i> , 2003, 184, 187-200.	1.9	21
18	Modelling, Steady State Analysis and Optimization of the Catalytic Efficiency of the Triosephosphate Isomerase. <i>Bulletin of Mathematical Biology</i> , 2002, 64, 301-326.	1.9	13

#	ARTICLE	IF	CITATIONS
19	Optimization of Tryptophan Production in Bacteria. Design of a Strategy for Genetic Manipulation of the Tryptophan Operon for Tryptophan Flux Maximization. Biotechnology Progress, 2000, 16, 133-145.	2.6	46