Markus Krummenacker

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3614858/publications.pdf

Version: 2024-02-01

20 papers 5,161 citations

16 h-index 794594 19 g-index

20 all docs

20 docs citations

times ranked

20

9178 citing authors

#	Article	IF	CITATIONS
1	Pathway Tools Management of Pathway/Genome Data for Microbial Communities. Frontiers in Bioinformatics, 2022, 2, .	2.1	4
2	Pathway Tools version 23.0 update: software for pathway/genome informatics and systems biology. Briefings in Bioinformatics, 2021, 22, 109-126.	6.5	117
3	Pathway Tools Visualization of Organism-Scale Metabolic Networks. Metabolites, 2021, 11, 64.	2.9	9
4	The EcoCyc Database in 2021. Frontiers in Microbiology, 2021, 12, 711077.	3.5	122
5	Metabolic route computation in organism communities. Microbiome, 2019, 7, 89.	11.1	4
6	The BioCyc collection of microbial genomes and metabolic pathways. Briefings in Bioinformatics, 2019, 20, 1085-1093.	6.5	582
7	The MetaCyc database of metabolic pathways and enzymes. Nucleic Acids Research, 2018, 46, D633-D639.	14.5	658
8	Pathway Tools version 19.0 update: software for pathway/genome informatics and systems biology. Briefings in Bioinformatics, 2016, 17, 877-890.	6.5	250
9	Computational Metabolomics Operations at BioCyc.org. Metabolites, 2015, 5, 291-310.	2.9	26
10	The MetaCyc database of metabolic pathways and enzymes and the BioCyc collection of Pathway/Genome Databases. Nucleic Acids Research, 2014, 42, D459-D471.	14.5	1,023
11	Optimal metabolic route search based on atom mappings. Bioinformatics, 2014, 30, 2043-2050.	4.1	47
12			
12	EcoCyc: fusing model organism databases with systems biology. Nucleic Acids Research, 2013, 41, D605-D612.	14.5	505
13	EcoCyc: fusing model organism databases with systems biology. Nucleic Acids Research, 2013, 41, D605-D612. Construction and completion of flux balance models from pathway databases. Bioinformatics, 2012, 28, 388-396.	14.5	505
	D605-D612. Construction and completion of flux balance models from pathway databases. Bioinformatics, 2012,		
13	D605-D612. Construction and completion of flux balance models from pathway databases. Bioinformatics, 2012, 28, 388-396. Pathway Tools version 13.0: integrated software for pathway/genome informatics and systems biology.	4.1	86
13	Construction and completion of flux balance models from pathway databases. Bioinformatics, 2012, 28, 388-396. Pathway Tools version 13.0: integrated software for pathway/genome informatics and systems biology. Briefings in Bioinformatics, 2010, 11, 40-79. Creating Fungal Pathway/Genome Databases Using Pathway Tools. Applied Mycology and	4.1 6.5	86 551
13 14 15	Construction and completion of flux balance models from pathway databases. Bioinformatics, 2012, 28, 388-396. Pathway Tools version 13.0: integrated software for pathway/genome informatics and systems biology. Briefings in Bioinformatics, 2010, 11, 40-79. Creating Fungal Pathway/Genome Databases Using Pathway Tools. Applied Mycology and Biotechnology, 2006, 6, 209-225. MetaCyc: a multiorganism database of metabolic pathways and enzymes. Nucleic Acids Research, 2006,	4.1 6.5 0.3	86 551 0

#	Article	IF	CITATIONS
19	Integrated pathway–genome databases and their role in drug discovery. Trends in Biotechnology, 1999, 17, 275-281.	9.3	154
20	EcoCyc: Enyclopedia of Escherichia coli Genes and Metabolism. Nucleic Acids Research, 1997, 25, 43-50.	14.5	46