Ron Caspi

List of Publications by Year in descending order

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279798 477307 9,124 29 23 29 citations h-index g-index papers 29 29 29 12871 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The MetaCyc database of metabolic pathways and enzymes and the BioCyc collection of pathway/genome databases. Nucleic Acids Research, 2016, 44, D471-D480.	14.5	1,788
2	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
3	The MetaCyc database of metabolic pathways and enzymes. Nucleic Acids Research, 2018, 46, D633-D639.	14.5	658
4	The MetaCyc database of metabolic pathways and enzymes - a 2019 update. Nucleic Acids Research, 2020, 48, D445-D453.	14.5	606
5	The MetaCyc Database of metabolic pathways and enzymes and the BioCyc collection of Pathway/Genome Databases. Nucleic Acids Research, 2007, 36, D623-D631.	14.5	600
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 and the BioCyc collection of pathway/genome databases. Nucleic Acids Research, 2012, 40, D742-D753.	14.5	561
8	Pathway Tools version 13.0: integrated software for pathway/genome informatics and systems biology. Briefings in Bioinformatics, 2010, 11, 40-79.	6.5	551
9	The EcoCyc database: reflecting new knowledge about <i>Escherichia coli</i> K-12. Nucleic Acids Research, 2017, 45, D543-D550.	14.5	541
10	MetaCyc: a multiorganism database of metabolic pathways and enzymes. Nucleic Acids Research, 2006, 34, D511-D516.	14.5	436
11	The MetaCyc database of metabolic pathways and enzymes and the BioCyc collection of pathway/genome databases. Nucleic Acids Research, 2010, 38, D473-D479.	14.5	403
12	Pathway Tools version 19.0 update: software for pathway/genome informatics and systems biology. Briefings in Bioinformatics, 2016, 17, 877-890.	6.5	250
13	Pathway size matters: the influence of pathway granularity on over-representation (enrichment) Tj ETQq $1\ 1\ 0.78$	4314 rgBT	 Overlock 10 -
14	A systematic comparison of the MetaCyc and KEGG pathway databases. BMC Bioinformatics, 2013, 14, 112.	2.6	123
15	The EcoCyc Database in 2021. Frontiers in Microbiology, 2021, 12, 711077.	3.5	122
16	Pathway Tools version 23.0 update: software for pathway/genome informatics and systems biology. Briefings in Bioinformatics, 2021, 22, 109-126.	6.5	117
17	Creation of a Genome-Wide Metabolic Pathway Database for <i>Populus trichocarpa</i> Using a New Approach for Reconstruction and Curation of Metabolic Pathways for Plants Â. Plant Physiology, 2010, 153, 1479-1491.	4.8	115
18	The Pathway Tools Pathway Prediction Algorithm. Standards in Genomic Sciences, 2011, 5, 424-429.	1.5	109

#	Article	IF	CITATIONS
19	The challenge of constructing, classifying, and representing metabolic pathways. FEMS Microbiology Letters, 2013, 345, 85-93.	1.8	82
20	The EcoCyc Database. EcoSal Plus, 2018, 8, .	5 . 4	75
21	A survey of metabolic databases emphasizing the MetaCyc family. Archives of Toxicology, 2011, 85, 1015-1033.	4.2	72
22	Sequence Analysis of a 101-Kilobase Plasmid Required for Agar Degradation by a Microscilla Isolate. Applied and Environmental Microbiology, 2001, 67, 5771-5779.	3.1	52
23	Beyond the genome (BTG) is a (PGDB) pathway genome database: HumanCyc. Genome Biology, 2010, 11, O12.	9.6	28
24	Nucleotide sequence based characterizations of two cryptic plasmids from the marine bacterium Ruegeria isolate PR1b. Plasmid, 2003, 49, 233-252.	1.4	25
25	Using the MetaCyc Pathway Database and the BioCyc Database Collection. Current Protocols in Bioinformatics, 2007, 20, Unit1.17.	25.8	25
26	Detection of DNA Replication Intermediates after Two-Dimensional Agarose Gel Electrophoresis Using a Fluorescein-Labeled Probe. Analytical Biochemistry, 1999, 269, 221-222.	2.4	11
27	A 50-kb Plasmid Rich in Mobile Gene Sequences Isolated from a Marine Micrococcus. Plasmid, 2002, 47, 1-9.	1.4	8
28	Plant Metabolic Pathways in MetaCyc and SolCyc. Nature Precedings, 2009, , .	0.1	2
29	Prediction of Selected Biosynthetic Pathways for the Lipopolysaccharide Components in Porphyromonas gingivalis. Pathogens, 2021, 10, 374.	2.8	1