

Amanda Clare

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1588277/publications.pdf>

Version: 2024-02-01

23
papers

968
citations

932766
10
h-index

794141
19
g-index

33
all docs

33
docs citations

33
times ranked

1510
citing authors

#	ARTICLE	IF	CITATIONS
1	The Automation of Science. <i>Science</i> , 2009, 324, 85-89.	6.0	458
2	Measuring scientific impact beyond academia: An assessment of existing impact metrics and proposed improvements. <i>PLoS ONE</i> , 2017, 12, e0173152.	1.1	118
3	Towards Robot Scientists for autonomous scientific discovery. <i>Automated Experimentation</i> , 2010, 2, 1.	2.0	101
4	An ontology for a Robot Scientist. <i>Bioinformatics</i> , 2006, 22, e464-e471.	1.8	56
5	Accurate Prediction of Protein Functional Class from Sequence in the <i>Mycobacterium tuberculosis</i> and <i>Escherichia coli</i> Genomes Using Data Mining. <i>Yeast</i> , 2000, 1, 283-293.	0.8	42
6	The EXACT description of biomedical protocols. <i>Bioinformatics</i> , 2008, 24, i295-i303.	1.8	31
7	The Robot Scientist Adam. <i>Computer</i> , 2009, 42, 46-54.	1.2	28
8	Replicating complex agent based models, a formidable task. <i>Environmental Modelling and Software</i> , 2017, 92, 142-151.	1.9	22
9	How well do we understand the clusters found in microarray data?. <i>In Silico Biology</i> , 2002, 2, 511-22.	0.4	22
10	No one tool to rule them all: prokaryotic gene prediction tool annotations are highly dependent on the organism of study. <i>Bioinformatics</i> , 2022, 38, 1198-1207.	1.8	20
11	On the complexity of haplotyping a microbial community. <i>Bioinformatics</i> , 2021, 37, 1360-1366.	1.8	17
12	AutoLabDB: a substantial open source database schema to support a high-throughput automated laboratory. <i>Bioinformatics</i> , 2012, 28, 1390-1397.	1.8	8
13	A natural language system for retrieval of captioned images. <i>Natural Language Engineering</i> , 2001, 7, 117-142.	2.1	7
14	PD5: A General Purpose Library for Primer Design Software. <i>PLoS ONE</i> , 2013, 8, e80156.	1.1	5
15	Evolutionary search techniques for the Lyndon factorization of biosequences. , 2019, , .		4
16	Wiki based management of chemometric research projects. <i>Journal of Chemometrics</i> , 2010, 24, 408-417.	0.7	3
17	Goldilocks: a tool for identifying genomic regions that are "just right". <i>Bioinformatics</i> , 2016, 32, 2047-2049.	1.8	3
18	Enhanced string factoring from alphabet orderings. <i>Information Processing Letters</i> , 2019, 143, 4-7.	0.4	2

#	ARTICLE	IF	CITATIONS
19	A Tool for Multiple Targeted Genome Deletions that Is Precise, Scar-Free, and Suitable for Automation. PLoS ONE, 2015, 10, e0142494.	1.1	1
20	Inductive Queries for a Drug Designing Robot Scientist. , 2010, , 425-451.		1
21	Laboratory Automation in a Functional Programming Language. Journal of the Association for Laboratory Automation, 2014, 19, 569-576.	2.8	0
22	Review of "A functional start to computing with Python", Ted Herman, CRC Press, 2014, ISBN 978-1-4665-0455-4. Journal of Functional Programming, 2015, 25, .	0.5	0
23	Evaluation of a Permutation-Based Evolutionary Framework for Lyndon Factorizations. Lecture Notes in Computer Science, 2020, , 390-403.	1.0	0