

Dave Clements

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

6,444
citations

567144

15
h-index

752573

20
g-index

27
all docs

27
docs citations

27
times ranked

12898
citing authors

#	ARTICLE	IF	CITATIONS
1	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2018 update. <i>Nucleic Acids Research</i> , 2018, 46, W537-W544.	6.5	3,003
2	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2016 update. <i>Nucleic Acids Research</i> , 2016, 44, W3-W10.	6.5	1,751
3	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2022 update. <i>Nucleic Acids Research</i> , 2022, 50, W345-W351.	6.5	328
4	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2020 update. <i>Nucleic Acids Research</i> , 2020, 48, W395-W402.	6.5	322
5	The Zebrafish Information Network: the zebrafish model organism database. <i>Nucleic Acids Research</i> , 2006, 34, D581-D585.	6.5	263
6	Community-Driven Data Analysis Training for Biology. <i>Cell Systems</i> , 2018, 6, 752-758.e1.	2.9	141
7	A high-resolution anatomical ontology of the developing murine genitourinary tract. <i>Gene Expression Patterns</i> , 2007, 7, 680-699.	0.3	125
8	Organization of the pronephric kidney revealed by large-scale gene expression mapping. <i>Genome Biology</i> , 2008, 9, R84.	13.9	116
9	The Zebrafish Information Network (ZFIN): the zebrafish model organism database. <i>Nucleic Acids Research</i> , 2003, 31, 241-243.	6.5	113
10	Inverting the model of genomics data sharing with the NHGRI Genomic Data Science Analysis, Visualization, and Informatics Lab-space. <i>Cell Genomics</i> , 2022, 2, 100085.	3.0	59
11	MetaBase--the wiki-database of biological databases. <i>Nucleic Acids Research</i> , 2012, 40, D1250-D1254.	6.5	40
12	Using Galaxy to Perform Large-Scale Interactive Data Analyses. <i>Current Protocols in Bioinformatics</i> , 2012, 38, Unit10.5.	25.8	36
13	The Chado Natural Diversity module: a new generic database schema for large-scale phenotyping and genotyping data. <i>Database: the Journal of Biological Databases and Curation</i> , 2011, 2011, bar051-bar051.	1.4	30
14	Myriad: Design and implementation of a federated database prototype. <i>Software - Practice and Experience</i> , 1995, 25, 533-562.	2.5	21
15	Fostering accessible online education using Galaxy as an e-learning platform. <i>PLoS Computational Biology</i> , 2021, 17, e1008923.	1.5	15
16	A single-cell RNA-sequencing training and analysis suite using the Galaxy framework. <i>GigaScience</i> , 2020, 9, .	3.3	14
17	Crowdsourcing biocuration: The Community Assessment of Community Annotation with Ontologies (CACAO). <i>PLoS Computational Biology</i> , 2021, 17, e1009463.	1.5	7
18	Using Galaxy to Perform Large-Scale Interactive Data Analyses--An Update. <i>Current Protocols</i> , 2021, 1, e31.	1.3	4

#	ARTICLE	IF	CITATIONS
19	The Galaxy Track Browser: Transforming the genome browser from visualization tool to analysis tool. , 2011, , .		1
20	Scaling Galaxy for Big Data. EMBnet Journal, 2014, 20, 764.	0.2	0