

# Dave Clements

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7167027/publications.pdf>

Version: 2024-02-01

20  
papers

6,444  
citations

567281

15  
h-index

752698

20  
g-index

27  
all docs

27  
docs citations

27  
times ranked

12898  
citing authors

#	ARTICLE	IF	CITATIONS
1	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.	6.5	59
2	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2022 update. <i>Nucleic Acids Research</i> , 2022, 50, W345-W351.	14.5	328
3	Using Galaxy to Perform Large-Scale Interactive Data Analyses—An Update. <i>Current Protocols</i> , 2021, 1, e31.	2.9	4
4	Fostering accessible online education using Galaxy as an e-learning platform. <i>PLoS Computational Biology</i> , 2021, 17, e1008923.	3.2	15
5	Crowdsourcing biocuration: The Community Assessment of Community Annotation with Ontologies (CACAO). <i>PLoS Computational Biology</i> , 2021, 17, e1009463.	3.2	7
6	A single-cell RNA-sequencing training and analysis suite using the Galaxy framework. <i>GigaScience</i> , 2020, 9, .	6.4	14
7	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2020 update. <i>Nucleic Acids Research</i> , 2020, 48, W395-W402.	14.5	322
8	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2018 update. <i>Nucleic Acids Research</i> , 2018, 46, W537-W544.	14.5	3,003
9	Community-Driven Data Analysis Training for Biology. <i>Cell Systems</i> , 2018, 6, 752-758.e1.	6.2	141
10	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2016 update. <i>Nucleic Acids Research</i> , 2016, 44, W3-W10.	14.5	1,751
11	Scaling Galaxy for Big Data. <i>EMBnet Journal</i> , 2014, 20, 764.	0.6	0
12	MetaBase--the wiki-database of biological databases. <i>Nucleic Acids Research</i> , 2012, 40, D1250-D1254.	14.5	40
13	Using Galaxy to Perform Large-Scale Interactive Data Analyses. <i>Current Protocols in Bioinformatics</i> , 2012, 38, Unit10.5.	25.8	36
14	The Galaxy Track Browser: Transforming the genome browser from visualization tool to analysis tool. , 2011, , .		1
15	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.	3.0	30
16	Organization of the pronephric kidney revealed by large-scale gene expression mapping. <i>Genome Biology</i> , 2008, 9, R84.	9.6	116
17	A high-resolution anatomical ontology of the developing murine genitourinary tract. <i>Gene Expression Patterns</i> , 2007, 7, 680-699.	0.8	125
18	The Zebrafish Information Network: the zebrafish model organism database. <i>Nucleic Acids Research</i> , 2006, 34, D581-D585.	14.5	263

#	ARTICLE	IF	CITATIONS
19	The Zebrafish Information Network (ZFIN): the zebrafish model organism database. <i>Nucleic Acids Research</i> , 2003, 31, 241-243.	14.5	113
20	Myriad: Design and implementation of a federated database prototype. <i>Software - Practice and Experience</i> , 1995, 25, 533-562.	3.6	21