

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

567144

15
h-index

752573

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. | 3.0 | 59 |
| 2 | The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2022 update. <i>Nucleic Acids Research</i> , 2022, 50, W345-W351. | 6.5 | 328 |
| 3 | Using Galaxy to Perform Large-Scale Interactive Data Analyses—An Update. <i>Current Protocols</i> , 2021, 1, e31. | 1.3 | 4 |
| 4 | Fostering accessible online education using Galaxy as an e-learning platform. <i>PLoS Computational Biology</i> , 2021, 17, e1008923. | 1.5 | 15 |
| 5 | Crowdsourcing biocuration: The Community Assessment of Community Annotation with Ontologies (CACAO). <i>PLoS Computational Biology</i> , 2021, 17, e1009463. | 1.5 | 7 |
| 6 | A single-cell RNA-sequencing training and analysis suite using the Galaxy framework. <i>GigaScience</i> , 2020, 9, . | 3.3 | 14 |
| 7 | The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2020 update. <i>Nucleic Acids Research</i> , 2020, 48, W395-W402. | 6.5 | 322 |
| 8 | 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 |
| 9 | Community-Driven Data Analysis Training for Biology. <i>Cell Systems</i> , 2018, 6, 752-758.e1. | 2.9 | 141 |
| 10 | 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 |
| 11 | Scaling Galaxy for Big Data. <i>EMBnet Journal</i> , 2014, 20, 764. | 0.2 | 0 |
| 12 | MetaBase--the wiki-database of biological databases. <i>Nucleic Acids Research</i> , 2012, 40, D1250-D1254. | 6.5 | 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. | 1.4 | 30 |
| 16 | Organization of the pronephric kidney revealed by large-scale gene expression mapping. <i>Genome Biology</i> , 2008, 9, R84. | 13.9 | 116 |
| 17 | A high-resolution anatomical ontology of the developing murine genitourinary tract. <i>Gene Expression Patterns</i> , 2007, 7, 680-699. | 0.3 | 125 |
| 18 | The Zebrafish Information Network: the zebrafish model organism database. <i>Nucleic Acids Research</i> , 2006, 34, D581-D585. | 6.5 | 263 |

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|----|---|-----|-----------|
| 19 | The Zebrafish Information Network (ZFIN): the zebrafish model organism database. <i>Nucleic Acids Research</i> , 2003, 31, 241-243. | 6.5 | 113 |
| 20 | Myriad: Design and implementation of a federated database prototype. <i>Software - Practice and Experience</i> , 1995, 25, 533-562. | 2.5 | 21 |