

Frederic B Bastian

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

631
citations

840119

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1125271

13
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16
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docs citations

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times ranked

1437
citing authors

#	ARTICLE	IF	CITATIONS
1	The Bgee suite: integrated curated expression atlas and comparative transcriptomics in animals. <i>Nucleic Acids Research</i> , 2021, 49, D831-D847.	6.5	112
2	OncoMX: A Knowledgebase for Exploring Cancer Biomarkers in the Context of Related Cancer and Healthy Data. <i>JCO Clinical Cancer Informatics</i> , 2020, 4, 210-220.	1.0	30
3	The SIB Swiss Institute of Bioinformatics's™ resources: focus on curated databases. <i>Nucleic Acids Research</i> , 2016, 44, D27-D37.	6.5	64
4	BgeeDB, an R package for retrieval of curated expression datasets and for gene list expression localization enrichment tests. <i>F1000Research</i> , 2016, 5, 2748.	0.8	23
5	BgeeDB, an R package for retrieval of curated expression datasets and for gene list expression localization enrichment tests. <i>F1000Research</i> , 2016, 5, 2748.	0.8	27
6	The Confidence Information Ontology: a step towards a standard for asserting confidence in annotations. <i>Database: the Journal of Biological Databases and Curation</i> , 2015, 2015, bav043-bav043.	1.4	37
7	Fifteen years SIB Swiss Institute of Bioinformatics: life science databases, tools and support. <i>Nucleic Acids Research</i> , 2014, 42, W436-W441.	6.5	13
8	Unification of multi-species vertebrate anatomy ontologies for comparative biology in Uberon. <i>Journal of Biomedical Semantics</i> , 2014, 5, 21.	0.9	121
9	Uncovering hidden duplicated content in public transcriptomics data. <i>Database: the Journal of Biological Databases and Curation</i> , 2013, 2013, bat010.	1.4	14
10	vHOG, a multispecies vertebrate ontology of homologous organs groups. <i>Bioinformatics</i> , 2012, 28, 1017-1020.	1.8	19
11	Homolonto: generating homology relationships by pairwise alignment of ontologies and application to vertebrate anatomy. <i>Bioinformatics</i> , 2010, 26, 1766-1771.	1.8	20
12	Bgee: Integrating and Comparing Heterogeneous Transcriptome Data Among Species. <i>Lecture Notes in Computer Science</i> , 2008, , 124-131.	1.0	137