

Pierre Larmande

List of Publications by Year in descending order

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Version: 2024-02-01

33
papers

1,418
citations

516710

16
h-index

414414

32
g-index

43
all docs

43
docs citations

43
times ranked

2386
citing authors

#	ARTICLE	IF	CITATIONS
1	High throughput T-DNA insertion mutagenesis in rice: a first step towards in silico reverse genetics. <i>Plant Journal</i> , 2004, 39, 450-464.	5.7	231
2	Highly efficient production and characterization of T-DNA plants for rice (<i>Oryza sativa</i> L.) functional genomics. <i>Theoretical and Applied Genetics</i> , 2003, 106, 1396-1408.	3.6	227
3	Scientific workflows for computational reproducibility in the life sciences: Status, challenges and opportunities. <i>Future Generation Computer Systems</i> , 2017, 75, 284-298.	7.5	104
4	A new cacao linkage map based on codominant markers: development and integration of 201 new microsatellite markers. <i>Theoretical and Applied Genetics</i> , 2004, 108, 1151-1161.	3.6	101
5	AgroPortal: A vocabulary and ontology repository for agronomy. <i>Computers and Electronics in Agriculture</i> , 2018, 144, 126-143.	7.7	87
6	The Rise and Fall of African Rice Cultivation Revealed by Analysis of 246 New Genomes. <i>Current Biology</i> , 2018, 28, 2274-2282.e6.	3.9	84
7	OryGenesDB: a database for rice reverse genetics. <i>Nucleic Acids Research</i> , 2006, 34, D736-D740.	14.5	82
8	BrAPI – an application programming interface for plant breeding applications. <i>Bioinformatics</i> , 2019, 35, 4147-4155.	4.1	82
9	P-TRAP: a Panicle Trait Phenotyping tool. <i>BMC Plant Biology</i> , 2013, 13, 122.	3.6	67
10	Oryza Tag Line , a phenotypic mutant database for the GÃ©noplante rice insertion line library. <i>Nucleic Acids Research</i> , 2008, 36, D1022-D1027.	14.5	60
11	AgBioData consortium recommendations for sustainable genomics and genetics databases for agriculture. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, .	3.0	52
12	OryGenesDB 2008 update: database interoperability for functional genomics of rice. <i>Nucleic Acids Research</i> , 2009, 37, D992-D995.	14.5	34
13	COMOKIT: A Modeling Kit to Understand, Analyze, and Compare the Impacts of Mitigation Policies Against the COVID-19 Epidemic at the Scale of a City. <i>Frontiers in Public Health</i> , 2020, 8, 563247.	2.7	34
14	In-depth molecular and phenotypic characterization in a rice insertion line library facilitates gene identification through reverse and forward genetics approaches. <i>Plant Biotechnology Journal</i> , 2012, 10, 555-568.	8.3	20
15	Gigwa – Genotype investigator for genome-wide analyses. <i>GigaScience</i> , 2016, 5, 25.	6.4	20
16	Gigwa v2 – Extended and improved genotype investigator. <i>GigaScience</i> , 2019, 8, .	6.4	20
17	Developing data interoperability using standards: A wheat community use case. <i>F1000Research</i> , 2017, 6, 1843.	1.6	20
18	Developing data interoperability using standards: A wheat community use case. <i>F1000Research</i> , 2017, 6, 1843.	1.6	14

#	ARTICLE	IF	CITATIONS
19	Agronomic Linked Data (AgroLD): A knowledge-based system to enable integrative biology in agronomy. PLoS ONE, 2018, 13, e0198270.	2.5	13
20	Rice Galaxy: an open resource for plant science. GigaScience, 2019, 8, .	6.4	11
21	Benchmarking database systems for Genomic Selection implementation. Database: the Journal of Biological Databases and Curation, 2019, 2019, .	3.0	7
22	Development of a knowledge system for Big Data. , 2016, , .		5
23	Evaluating Named-Entity Recognition Approaches in Plant Molecular Biology. Lecture Notes in Computer Science, 2018, , 219-225.	1.3	4
24	Progress in single-access information systems for wheat and rice crop improvement. Briefings in Bioinformatics, 2019, 20, 565-571.	6.5	4
25	AgroLD: A Knowledge Graph for the Plant Sciences. Lecture Notes in Computer Science, 2021, , 496-510.	1.3	3
26	OryzaGP: rice gene and protein dataset for named-entity recognition. Genomics and Informatics, 2019, 17, e17.	0.8	3
27	Clever generation of rich SPARQL queries from annotated relational schema: application to Semantic Web Service creation for biological databases. BMC Bioinformatics, 2013, 14, 126.	2.6	2
28	Enabling a fast annotation process with the Table2Annotation tool. Genomics and Informatics, 2020, 18, e19.	0.8	2
29	AgroLD: A Knowledge Graph Database for Plant Functional Genomics. Methods in Molecular Biology, 2022, 2443, 527-540.	0.9	2
30	OryzaGP 2021 update: a rice gene and protein dataset for named-entity recognition. Genomics and Informatics, 2021, 19, e27.	0.8	1
31	Managing High-Density Genotyping Data with Gigwa. Methods in Molecular Biology, 2022, 2443, 415-427.	0.9	1
32	PyRice: a Python package for querying <i>Oryza sativa</i> databases. Bioinformatics, 2021, 37, 1037-1038.	4.1	0
33	AgroLD API AgroLD API. Une architecture orientée services pour l'extraction de connaissances dans la base de données liées AgroLD. Ingenierie Des Systemes D'Information, 2016, 21, 133-157.	0.7	0