Samuel Lampa

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/833100/publications.pdf

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16	626	12 h-index	13
papers	citations		g-index
21	21	21	1377 citing authors
all docs	docs citations	times ranked	

#	Article	IF	Citations
1	Towards reproducible computational drug discovery. Journal of Cheminformatics, 2020, 12, 9.	2.8	100
2	Software engineering for scientific big data analysis. GigaScience, 2019, 8, .	3.3	20
3	SciPipe—Turning Scientific Workflows into Computer Programs. Computing in Science and Engineering, 2019, 21, 109-113.	1.2	O
4	SciPipe: A workflow library for agile development of complex and dynamic bioinformatics pipelines. GigaScience, 2019, 8, .	3.3	22
5	PhenoMeNal: processing and analysis of metabolomics data in the cloud. GigaScience, 2019, 8, .	3.3	60
6	Predicting Off-Target Binding Profiles With Confidence Using Conformal Prediction. Frontiers in Pharmacology, 2018, 9, 1256.	1.6	21
7	A confidence predictor for logD using conformal regression and a support-vector machine. Journal of Cheminformatics, 2018, 10, 17.	2.8	37
8	SweGen: a whole-genome data resource of genetic variability in a cross-section of the Swedish population. European Journal of Human Genetics, 2017, 25, 1253-1260.	1.4	148
9	RDFIO: extending Semantic MediaWiki for interoperable biomedical data management. Journal of Biomedical Semantics, 2017, 8, 35.	0.9	5
10	Towards agile large-scale predictive modelling in drug discovery with flow-based programming design principles. Journal of Cheminformatics, 2016, 8, 67.	2.8	26
11	Large-scale ligand-based predictive modelling using support vector machines. Journal of Cheminformatics, 2016, 8, 39.	2.8	34
12	Experiences with workflows for automating data-intensive bioinformatics. Biology Direct, 2015, 10, 43.	1.9	52
13	Lessons learned from implementing a national infrastructure in Sweden for storage and analysis of next-generation sequencing data. GigaScience, 2013, 2, 9.	3.3	67
14	UPPNEX - A solution for Next Generation Sequencing data management and analysis. EMBnet Journal, 2012, 17, 44.	0.2	0
15	Linking the Resource Description Framework to cheminformatics and proteochemometrics. Journal of Biomedical Semantics, 2011, 2, S6.	0.9	24
16	Approaches for containerized scientific workflows in cloud environments with applications in life science. F1000Research, 0, 10, 513.	0.8	1