

Chelsea J T Ju

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1760605/publications.pdf>

Version: 2024-02-01

10
papers

489
citations

1478505

6
h-index

1474206

9
g-index

14
all docs

14
docs citations

14
times ranked

953
citing authors

#	ARTICLE	IF	CITATIONS
1	Multifaceted proteinâ€“protein interaction prediction based on Siamese residual RCNN. <i>Bioinformatics</i> , 2019, 35, i305-i314.	4.1	181
2	Leveraging molecular quantitative trait loci to understand the genetic architecture of diseases and complex traits. <i>Nature Genetics</i> , 2018, 50, 1041-1047.	21.4	154
3	Widespread Allelic Heterogeneity in Complex Traits. <i>American Journal of Human Genetics</i> , 2017, 100, 789-802.	6.2	74
4	Mutation effect estimation on proteinâ€“protein interactions using deep contextualized representation learning. <i>NAR Genomics and Bioinformatics</i> , 2020, 2, lqaa015.	3.2	41
5	Functional disease architectures reveal unique biological role of transposable elements. <i>Nature Communications</i> , 2019, 10, 4054.	12.8	14
6	JEDI: circular RNA prediction based on junction encoders and deep interaction among splice sites. <i>Bioinformatics</i> , 2021, 37, i289-i298.	4.1	11
7	Prediction of microbial communities for urban metagenomics using neural network approach. <i>Human Genomics</i> , 2019, 13, 47.	2.9	6
8	Efficient Approach to Correct Read Alignment for Pseudogene Abundance Estimates. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2017, 14, 522-533.	3.0	4
9	TahcoRoll: fast genomic signature profiling via thinned automaton and rolling hash. <i>Medical Review</i> , 2022, .	1.2	2
10	Inferring Microbial Communities for City Scale Metagenomics Using Neural Networks. , 2018, , .		0