

Dan DeBlasio

List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

392
citations

1307594

7
h-index

888059

17
g-index

41
all docs

41
docs citations

41
times ranked

514
citing authors

#	ARTICLE	IF	CITATIONS
1	How much data is sufficient to learn high-performing algorithms? generalization guarantees for data-driven algorithm design. , 2021, , .		2
2	More Accurate Transcript Assembly via Parameter Advising. Journal of Computational Biology, 2020, 27, 1181-1189.	1.6	6
3	Locality-sensitive hashing for the edit distance. Bioinformatics, 2019, 35, i127-i135.	4.1	40
4	Practical Universal k-mer Sets for Minimizer Schemes. , 2019, , .		18
5	Nurturing tomorrow's leaders: The ISCB Student Council Symposia in 2018. F1000Research, 2019, 8, 34.	1.6	6
6	Global network of computational biology communities: ISCB's Regional Student Groups breaking barriers. F1000Research, 2019, 8, 1574.	1.6	8
7	Reflections on a journey: a retrospective of the ISCB Student Council symposium series. BMC Bioinformatics, 2018, 19, 347.	2.6	9
8	Asymptotically optimal minimizers schemes. Bioinformatics, 2018, 34, i13-i22.	4.1	44
9	Adaptive Local Realignment of Protein Sequences. Journal of Computational Biology, 2018, 25, 780-793.	1.6	1
10	The ISCB Student Council Internship Program: Expanding computational biology capacity worldwide. PLoS Computational Biology, 2018, 14, e1005802.	3.2	4
11	Learning Parameter-Advising Sets for Multiple Sequence Alignment. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2017, 14, 1028-1041.	3.0	10
12	Core column prediction for protein multiple sequence alignments. Algorithms for Molecular Biology, 2017, 12, 11.	1.2	2
13	Boosting Alignment Accuracy by Adaptive Local Realignment. Lecture Notes in Computer Science, 2017, , 1-17.	1.3	2
14	Parameter Advising for the Opal Aligner. Computational Biology, 2017, , 65-83.	0.2	2
15	Highlights of the second ISCB Student Council Symposium in Africa, 2017. F1000Research, 2017, 6, 2183.	1.6	6
16	Computational Complexity of Advising. Computational Biology, 2017, , 41-49.	0.2	0
17	Ensemble Multiple Alignment. Computational Biology, 2017, , 85-102.	0.2	0
18	Core Column Prediction for Alignments. Computational Biology, 2017, , 117-137.	0.2	0

#	ARTICLE	IF	CITATIONS
19	Alignment Accuracy Estimation. Computational Biology, 2017, , 19-27.	0.2	0
20	The Facet Accuracy Estimator. Computational Biology, 2017, , 29-40.	0.2	0
21	Constructing Advisors. Computational Biology, 2017, , 51-61.	0.2	0
22	Adaptive Local Realignment via Parameter Advising. , 2016, , .		0
23	Predicting Core Columns of Protein Multiple Sequence Alignments for Improved Parameter Advising. Lecture Notes in Computer Science, 2016, , 77-89.	1.3	2
24	Highlights from the 11th ISCB Student Council Symposium 2015. BMC Bioinformatics, 2016, 17, 95.	2.6	4
25	Parameter advising for multiple sequence alignment. BMC Bioinformatics, 2015, 16, .	2.6	8
26	Ensemble multiple sequence alignment via advising. , 2015, , .		8
27	Learning parameter sets for alignment advising. , 2014, , .		10
28	Accuracy Estimation and Parameter Advising for Protein Multiple Sequence Alignment. Journal of Computational Biology, 2013, 20, 259-279.	1.6	26
29	A Memory Efficient Method for Structure-Based RNA Multiple Alignment. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2012, 9, 1-11.	3.0	5
30	Epigenetic regulation of microRNA miR-375 and its role in melanoma development in humans. FEBS Letters, 2011, 585, 2467-2476.	2.8	85
31	Epigenetic Regulation of MicroRNA Genes and the Role of miR-34b in Cell Invasion and Motility in Human Melanoma. PLoS ONE, 2011, 6, e24922.	2.5	63