

# Alexandre P Francisco

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9504260/publications.pdf>

Version: 2024-02-01

15  
papers

1,831  
citations

933447

10  
h-index

1058476

14  
g-index

16  
all docs

16  
docs citations

16  
times ranked

2784  
citing authors

#	ARTICLE	IF	CITATIONS
1	VeilGraph: incremental graph stream processing. Journal of Big Data, 2022, 9, .	11.0	5
2	Shapelets to Classify Energy Demand Time Series. Energies, 2022, 15, 2960.	3.1	2
3	Distance-based phylogenetic inference from typing data: a unifying view. Briefings in Bioinformatics, 2021, 22, .	6.5	8
4	An analysis of the graph processing landscape. Journal of Big Data, 2021, 8, 55.	11.0	11
5	Eliciting Fairness in N-Player Network Games through Degree-Based Role Assignment. Complexity, 2021, 2021, 1-11.	1.6	5
6	On Dynamic Succinct Graph Representations. , 2020, , .		3
7	Using Machine Learning to Improve the Prediction of Functional Outcome in Ischemic Stroke Patients. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2018, 15, 1953-1959.	3.0	79
8	Fast phylogenetic inference from typing data. Algorithms for Molecular Biology, 2018, 13, 4.	1.2	14
9	PHYLOViZ 2.0: providing scalable data integration and visualization for multiple phylogenetic inference methods. Bioinformatics, 2017, 33, 128-129.	4.1	336
10	PHYLOViZ Online: web-based tool for visualization, phylogenetic inference, analysis and sharing of minimum spanning trees. Nucleic Acids Research, 2016, 44, W246-W251.	14.5	152
11	Extending quick hypervolume. Journal of Heuristics, 2016, 22, 245-271.	1.4	19
12	Not Seeing the Forest for the Trees: Size of the Minimum Spanning Trees (MSTs) Forest and Branch Significance in MST-Based Phylogenetic Analysis. PLoS ONE, 2015, 10, e0119315.	2.5	15
13	Quick Hypervolume. IEEE Transactions on Evolutionary Computation, 2014, 18, 481-502.	10.0	158
14	PHYLOViZ: phylogenetic inference and data visualization for sequence based typing methods. BMC Bioinformatics, 2012, 13, 87.	2.6	492
15	Global optimal eBURST analysis of multilocus typing data using a graphic matroid approach. BMC Bioinformatics, 2009, 10, 152.	2.6	522