## **Benoit Morel**

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/615633/publications.pdf

Version: 2024-02-01

1478505 1872680 2,745 6 6 6 citations h-index g-index papers 13 13 13 4541 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	SpeciesRax: A Tool for Maximum Likelihood Species Tree Inference from Gene Family Trees under Duplication, Transfer, and Loss. Molecular Biology and Evolution, 2022, 39, .	8.9	29
2	NetRAX: accurate and fast maximum likelihood phylogenetic network inference. Bioinformatics, 2022, 38, 3725-3733.	4.1	13
3	Phylogenetic Analysis of SARS-CoV-2 Data Is Difficult. Molecular Biology and Evolution, 2021, 38, 1777-1791.	8.9	119
4	GeneRax: A Tool for Species-Tree-Aware Maximum Likelihood-Based Gene ÂFamily Tree Inference under Gene Duplication, Transfer, and Loss. Molecular Biology and Evolution, 2020, 37, 2763-2774.	8.9	87
5	RAxML-NG: a fast, scalable and user-friendly tool for maximum likelihood phylogenetic inference. Bioinformatics, 2019, 35, 4453-4455.	4.1	2,287
6	ParGenes: a tool for massively parallel model selection and phylogenetic tree inference on thousands of genes. Bioinformatics, 2019, 35, 1771-1773.	4.1	51