

Claudia Solis-Lemus

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7080284/publications.pdf>

Version: 2024-02-01

15
papers

1,064
citations

1162367

8
h-index

1199166

12
g-index

18
all docs

18
docs citations

18
times ranked

1387
citing authors

#	ARTICLE	IF	CITATIONS
1	Efficient estimation of indirect effects in case-control studies using a unified likelihood framework. <i>Statistics in Medicine</i> , 2022, 41, 2879-2893.	0.8	2
2	Towards a robust out-of-the-box neural network model for genomic data. <i>BMC Bioinformatics</i> , 2022, 23, 125.	1.2	1
3	WI Fast Stats: a collection of web apps for the visualization and analysis of WI Fast Plants data. <i>The Journal of Open Source Education</i> , 2022, 5, 159.	0.2	0
4	Genes Influencing Phage Host Range in <i>Staphylococcus aureus</i> on a Species-Wide Scale. <i>MSphere</i> , 2021, 6, .	1.3	19
5	Effect of genetic background on the evolution of Vancomycin-Intermediate <i>Staphylococcus aureus</i> (VISA). <i>PeerJ</i> , 2021, 9, e11764.	0.9	9
6	Leveraging Family History in Case-Control Analyses of Rare Variation. <i>Genetics</i> , 2020, 214, 295-303.	1.2	0
7	Genomic analysis of variability in Delta-toxin levels between <i>Staphylococcus aureus</i> strains. <i>PeerJ</i> , 2020, 8, e8717.	0.9	12
8	A generalized measure of dispersion. <i>Statistics and Probability Letters</i> , 2020, 164, 108806.	0.4	0
9	Phylogenetic Comparative Methods on Phylogenetic Networks with Reticulations. <i>Systematic Biology</i> , 2018, 67, 800-820.	2.7	88
10	Greatly reduced phylogenetic structure in the cultivated potato clade (<i>Solanum</i> section) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 38	0.8	16
11	PhyloNetworks: A Package for Phylogenetic Networks. <i>Molecular Biology and Evolution</i> , 2017, 34, 3292-3298.	3.5	250
12	Inferring Phylogenetic Networks with Maximum Pseudolikelihood under Incomplete Lineage Sorting. <i>PLoS Genetics</i> , 2016, 12, e1005896.	1.5	348
13	Statistical evidence for common ancestry: Application to primates. <i>Evolution; International Journal of Organic Evolution</i> , 2016, 70, 1354-1363.	1.1	8
14	Inconsistency of Species Tree Methods under Gene Flow. <i>Systematic Biology</i> , 2016, 65, 843-851.	2.7	146
15	Bayesian species delimitation combining multiple genes and traits in a unified framework. <i>Evolution; International Journal of Organic Evolution</i> , 2015, 69, 492-507.	1.1	152