

Frederick J Boehm

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3182711/publications.pdf>

Version: 2024-02-01

10
papers

461
citations

1307594

7
h-index

1474206

9
g-index

14
all docs

14
docs citations

14
times ranked

612
citing authors

#	ARTICLE	IF	CITATIONS
1	Host-pathogen genetic interactions underlie tuberculosis susceptibility in genetically diverse mice. <i>ELife</i> , 2022, 11, .	6.0	44
2	Statistical methods for Mendelian randomization in genome-wide association studies: A review. <i>Computational and Structural Biotechnology Journal</i> , 2022, 20, 2338-2351.	4.1	56
3	What Is Happening on Twitter? A Framework for Student Research Projects With Tweets. <i>Journal of Statistics and Data Science Education</i> , 2021, 29, S95-S102.	1.6	3
4	Genetic background modifies phenotypic severity and longevity in a mouse model of Niemann-Pick disease type C1. <i>DMM Disease Models and Mechanisms</i> , 2020, 13, .	2.4	17
5	Genetic determinants of gut microbiota composition and bile acid profiles in mice. <i>PLoS Genetics</i> , 2019, 15, e1008073.	3.5	75
6	Testing Pleiotropy vs. Separate QTL in Multiparental Populations. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 2317-2324.	1.8	11
7	qtl2pleio: Testing pleiotropy vs. separate QTL in multiparental populations. <i>Journal of Open Source Software</i> , 2019, 4, 1435.	4.6	0
8	How Trump Drove Coverage to the Nomination: Hybrid Media Campaigning. <i>Political Communication</i> , 2016, 33, 669-676.	3.9	151
9	Stable kinetochore-microtubule attachment is sufficient to silence the spindle assembly checkpoint in human cells. <i>Nature Communications</i> , 2015, 6, 10036.	12.8	91
10	A Strategy to Identify Dominant Point Mutant Modifiers of a Quantitative Trait. <i>G3: Genes, Genomes, Genetics</i> , 2014, 4, 1113-1121.	1.8	4