## C Brandon Ogbunugafor

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3152796/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Exploring the expanse between theoretical questions and experimental approaches in the modern study of evolvability. Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2023, 340, 8-17.	0.6	2
2	The mutation effect reaction norm (muâ€rn) highlights environmentally dependent mutation effects and epistatic interactions. Evolution; International Journal of Organic Evolution, 2022, 76, 37-48.	1.1	12
3	The endoplasmic reticulum proteostasis network profoundly shapes the protein sequence space accessible to HIV envelope. PLoS Biology, 2022, 20, e3001569.	2.6	7
4	Molecular mechanisms and drivers of pathogen emergence. Trends in Microbiology, 2022, 30, 898-911.	3.5	19
5	Higher-Order Interactions in Biology: The Curious Case of Epistasis. Understanding Complex Systems, 2022, , 417-433.	0.3	3
6	Cholera dynamics: lessons from an epidemic. Journal of Medical Microbiology, 2021, 70, .	0.7	14
7	Variation in microparasite free-living survival and indirect transmission can modulate the intensity of emerging outbreaks. Scientific Reports, 2020, 10, 20786.	1.6	7
8	Direct transmission via households informs models of disease and intervention dynamics in cholera. PLoS ONE, 2020, 15, e0229837.	1.1	14
9	A Reflection on 50 Years of John Maynard Smith's "Protein Spaceâ€: Genetics, 2020, 214, 749-754.	1.2	13
10	Experimental evolution for niche breadth in bacteriophage T4 highlights the importance of structural genes. MicrobiologyOpen, 2020, 9, e968.	1.2	2
11	The Epidemiological Signature of Pathogen Populations That Vary in the Relationship between Free-Living Parasite Survival and Virulence. Viruses, 2020, 12, 1055.	1.5	5
12	Genetic Background Modifies the Topography of a Fitness Landscape, Influencing the Dynamics of Adaptive Evolution. IEEE Access, 2019, 7, 113675-113683.	2.6	6
13	Hepatitis C virus modelled as an indirectly transmitted infection highlights the centrality of injection drug equipment in disease dynamics. Journal of the Royal Society Interface, 2019, 16, 20190334.	1.5	9
14	Lexical Landscapes as large in silico data for examining advanced properties of fitness landscapes. PLoS ONE, 2019, 14, e0220891.	1.1	6
15	Chimeric dihydrofolate reductases display properties of modularity and biophysical diversity. Protein Science, 2019, 28, 1359-1367.	3.1	3
16	Proteostasis Environment Shapes Higher-Order Epistasis Operating on Antibiotic Resistance. Genetics, 2019, 212, 565-575.	1.2	30
17	Competition along trajectories governs adaptation rates towards antimicrobial resistance. Nature Ecology and Evolution, 2017, 1, 7.	3.4	26

18 Quantifying Deception. , 2016, , .

#	Article	IF	CITATIONS
19	A pivot mutation impedes reverse evolution across an adaptive landscape for drug resistance in Plasmodium vivax. Malaria Journal, 2016, 15, 40.	0.8	22
20	Adaptive Landscape by Environment Interactions Dictate Evolutionary Dynamics in Models of Drug Resistance. PLoS Computational Biology, 2016, 12, e1004710.	1.5	71
21	A New Take on John Maynard Smith's Concept of Protein Space for Understanding Molecular Evolution. PLoS Computational Biology, 2016, 12, e1005046.	1.5	5
22	OFFI Models: Novel Schema for Dynamical Modeling of Biological Systems. PLoS ONE, 2016, 11, e0156844.	1.1	2
23	Delayed transmission selects for increased survival of vesicular stomatitis virus. Evolution; International Journal of Organic Evolution, 2015, 69, 117-125.	1.1	24
24	Evolution of Increased Survival in RNA Viruses Specialized on Cancer-Derived Cells. American Naturalist, 2013, 181, 585-595.	1.0	13
25	Combining mathematics and empirical data to predict emergence of RNA viruses that differ in reservoir use. Philosophical Transactions of the Royal Society B: Biological Sciences, 2010, 365, 1919-1930.	1.8	8
26	On the possible role of robustness in the evolution of infectious diseases. Chaos, 2010, 20, 026108.	1.0	9