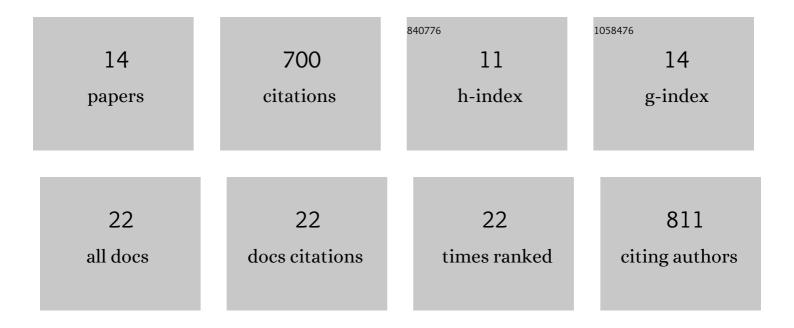
Ellie E Armstrong

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1008732/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The crucial role of genome-wide genetic variation in conservation. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	196
2	Widespread introgression across a phylogeny of 155 Drosophila genomes. Current Biology, 2022, 32, 111-123.e5.	3.9	132
3	Highly contiguous assemblies of 101 drosophilid genomes. ELife, 2021, 10, .	6.0	108
4	The Mutationathon highlights the importance of reaching standardization in estimates of pedigree-based germline mutation rates. ELife, 2022, 11, .	6.0	38
5	Horizontal Transfer of Bacterial Cytolethal Distending Toxin B Genes to Insects. Molecular Biology and Evolution, 2019, 36, 2105-2110.	8.9	36
6	Recent Evolutionary History of Tigers Highlights Contrasting Roles of Genetic Drift and Selection. Molecular Biology and Evolution, 2021, 38, 2366-2379.	8.9	35
7	Long live the king: chromosome-level assembly of the lion (Panthera leo) using linked-read, Hi-C, and long-read data. BMC Biology, 2020, 18, 3.	3.8	34
8	Rapid divergence of mussel populations despite incomplete barriers to dispersal. Molecular Ecology, 2018, 27, 1556-1571.	3.9	29
9	Comparative analyses identify genomic features potentially involved in the evolution of birds-of-paradise. GigaScience, 2019, 8, .	6.4	22
10	Cost-effective assembly of the African wild dog (<i>Lycaon pictus</i>) genome using linked reads. GigaScience, 2019, 8, .	6.4	22
11	The <i>Tetragnatha kauaiensis</i> Genome Sheds Light on the Origins of Genomic Novelty in Spiders. Genome Biology and Evolution, 2021, 13, .	2.5	16
12	Draft Genome Sequence and Annotation of the Lichen-Forming Fungus Arthonia radiata. Genome Announcements, 2018, 6, .	0.8	12
13	A holobiont view of island biogeography: Unravelling patterns driving the nascent diversification of a Hawaiian spider and its microbial associates. Molecular Ecology, 2022, 31, 1299-1316.	3.9	5
14	Good from far, but far from good: The impact of a reference genome on evolutionary inference. Molecular Ecology Resources, 2022, 22, 12-14.	4.8	2