

Holly L Lutz

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/695928/publications.pdf>

Version: 2024-02-01

24
papers

983
citations

567281

15
h-index

642732

23
g-index

31
all docs

31
docs citations

31
times ranked

1251
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative Analyses of Vertebrate Gut Microbiomes Reveal Convergence between Birds and Bats. MBio, 2020, 11, .	4.1	204
2	Climate variation influences host specificity in avian malaria parasites. Ecology Letters, 2019, 22, 547-557.	6.4	90
3	Parasite Prevalence Corresponds to Host Life History in a Diverse Assemblage of Afrotropical Birds and Haemosporidian Parasites. PLoS ONE, 2015, 10, e0121254.	2.5	87
4	Deeply conserved susceptibility in a multi-host, multi-parasite system. Ecology Letters, 2019, 22, 987-998.	6.4	54
5	Avian host composition, local speciation and dispersal drive the regional assembly of avian malaria parasites in South American birds. Molecular Ecology, 2019, 28, 2681-2693.	3.9	54
6	Ecology and Host Identity Outweigh Evolutionary History in Shaping the Bat Microbiome. MSystems, 2019, 4, .	3.8	54
7	An inverse latitudinal gradient in infection probability and phylogenetic diversity for <i>Leucocytozoon</i> blood parasites in New World birds. Journal of Animal Ecology, 2020, 89, 423-435.	2.8	49
8	Gut microbiota-driven brain A β amyloidosis in mice requires microglia. Journal of Experimental Medicine, 2022, 219, .	8.5	44
9	Diverse sampling of East African haemosporidians reveals chiropteran origin of malaria parasites in primates and rodents. Molecular Phylogenetics and Evolution, 2016, 99, 7-15.	2.7	43
10	Global drivers of avian haemosporidian infections vary across zoogeographical regions. Global Ecology and Biogeography, 2021, 30, 2393-2406.	5.8	42
11	Building Natural History Collections for the Twenty-First Century and Beyond. BioScience, 2020, 70, 674-687.	4.9	40
12	Leveraging natural history biorepositories as a global, decentralized, pathogen surveillance network. PLoS Pathogens, 2021, 17, e1009583.	4.7	38
13	Biogeography and spatio-temporal diversification of Selenidera and Andigena Toucans (Aves: Tj ETQq1 1 0.784314 rgBT / Overlock 10 27	2.7	27
14	Effects of Extended Postmortem Interval on Microbial Communities in Organs of the Human Cadaver. Frontiers in Microbiology, 2020, 11, 569630.	3.5	26
15	Laser capture microdissection microscopy and genome sequencing of the avian malaria parasite, Plasmodium relictum. Parasitology Research, 2016, 115, 4503-4510.	1.6	20
16	Multilocus phylogeny of a cryptic radiation of Afrotropical long-fingered bats (Chiroptera,) Tj ETQq0 0 0 rgBT / Overlock 10 Tj 50 142 T	1.7	17
17	Diversity of Sea Star-Associated Densoviruses and Transcribed Endogenous Viral Elements of Densovirus Origin. Journal of Virology, 2020, 95, .	3.4	14
18	A Simple Microbiome in the European Common Cuttlefish, <i>Sepia officinalis</i> . MSystems, 2019, 4, .	3.8	13

#	ARTICLE	IF	CITATIONS
19	Conceptual strategies for characterizing interactions in microbial communities. IScience, 2022, 25, 103775.	4.1	12
20	Curatorial guidelines and standards of the American Society of Mammalogists for collections of genetic resources. Journal of Mammalogy, 2019, 100, 1690-1694.	1.3	11
21	Associations between Afrotropical bats, eukaryotic parasites, and microbial symbionts. Molecular Ecology, 2022, 31, 1939-1950.	3.9	10
22	Microbiome is not linked to clinical disease severity of familial Mediterranean fever in an international cohort of children. Clinical and Experimental Rheumatology, 2021, 39, 102-108.	0.8	3
23	Wildlife Microbial Genomics and Endocrinology. , 2019, , .		0
24	Microbiome is not linked to clinical disease severity of familial Mediterranean fever in an international cohort of children. Clinical and Experimental Rheumatology, 2021, 39 Suppl 132, 102-108.	0.8	0