Elin Videvall

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7180402/publications.pdf

Version: 2024-02-01

24	837	15	22
papers	citations	h-index	g-index
33	33	33	1437
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Sharing and reporting benefits from biodiversity research. Molecular Ecology, 2021, 30, 1103-1107.	3.9	19
2	The uropygial gland microbiome of house sparrows with malaria infection. Journal of Avian Biology, 2021, 52, .	1.2	11
3	Transcriptome assembly and differential gene expression of the invasive avian malaria parasite <i>Plasmodium relictum</i> in Hawaiʻi. Ecology and Evolution, 2021, 11, 4935-4944.	1.9	10
4	Natural experiments and long-term monitoring are critical to understand and predict marine host–microbe ecology and evolution. PLoS Biology, 2021, 19, e3001322.	5 . 6	17
5	Early-life gut dysbiosis linked to juvenile mortality in ostriches. Microbiome, 2020, 8, 147.	11.1	30
6	Host Transcriptional Responses to High- and Low-Virulent Avian Malaria Parasites. American Naturalist, 2020, 195, 1070-1084.	2.1	19
7	Microbiome maturation during a unique developmental window. Molecular Ecology, 2020, 29, 1941-1943.	3.9	1
8	Major shifts in gut microbiota during development and its relationship to growth in ostriches. Molecular Ecology, 2019, 28, 2653-2667.	3.9	53
9	Genomic Advances in Avian Malaria Research. Trends in Parasitology, 2019, 35, 254-266.	3.3	23
10	Genomics of host-pathogen interactions: challenges and opportunities across ecological and spatiotemporal scales. PeerJ, 2019, 7, e8013.	2.0	23
11	De novo synthesis of thiamine (vitamin B1) is the ancestral state in Plasmodium parasites – evidence from avian haemosporidians. Parasitology, 2018, 145, 1084-1089.	1.5	2
12	Measuring the gut microbiome in birds: Comparison of faecal and cloacal sampling. Molecular Ecology Resources, 2018, 18, 424-434.	4.8	125
13	Insights into Avian Incomplete Dosage Compensation: Sex-Biased Gene Expression Coevolves with Sex Chromosome Degeneration in the Common Whitethroat. Genes, 2018, 9, 373.	2.4	13
14	Plasmodium parasites of birds have the most AT-rich genes of eukaryotes. Microbial Genomics, 2018, 4, .	2.0	25
15	The transcriptome of the avian malaria parasite <i>Plasmodium ashfordi</i> displays hostâ€specific gene expression. Molecular Ecology, 2017, 26, 2939-2958.	3.9	41
16	Transcriptome analysis of a wild bird reveals physiological responses to the urban environment. Scientific Reports, 2017, 7, 44180.	3.3	86
17	Direct PCR Offers a Fast and Reliable Alternative to Conventional DNA Isolation Methods for Gut Microbiomes. MSystems, 2017, 2, .	3.8	26
18	The Genome of <i>Haemoproteus tartakovskyi </i> and Its Relationship to Human Malaria Parasites. Genome Biology and Evolution, 2016, 8, 1361-1373.	2.5	58

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#	Article	IF	CITATION
19	Strong Maternal Effects on Gene Expression in <i> Arabidopsis lyrata < /i > Hybrids. Molecular Biology and Evolution, 2016, 33, 984-994.</i>	8.9	22
20	The Avian Transcriptome Response to Malaria Infection. Molecular Biology and Evolution, 2015, 32, 1255-1267.	8.9	83
21	Sex- and tissue-specific profiles of chemosensory gene expression in a herbivorous gall-inducing fly (Diptera: Cecidomyiidae). BMC Genomics, 2014, 15, 501.	2.8	81
22	Molecular identification of bloodmeals and species composition in <i>Culicoides</i> biting midges. Medical and Veterinary Entomology, 2013, 27, 104-112.	1.5	51
23	Butterfly monitoring using systematically placed transects in contrasting climatic regions – exploring an established spatial design for sampling. Nature Conservation, 0, 14, 41-62.	0.0	7
24	Personalized microbiomes in social baboons. Nature Ecology and Evolution, 0, , .	7.8	0