

Sara Jane Oppenheim

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

219
citations

1478280

6
h-index

1199470

12
g-index

15
all docs

15
docs citations

15
times ranked

344
citing authors

#	ARTICLE	IF	CITATIONS
1	We can't all be supermodels: the value of comparative transcriptomics to the study of non-model insects. <i>Insect Molecular Biology</i> , 2015, 24, 139-154.	1.0	82
2	BEHAVIORAL ADAPTATIONS INCREASE THE VALUE OF ENEMY-FREE SPACE FOR <i>HELIOTHIS SUBFLEXA</i> , A SPECIALIST HERBIVORE. <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 679-689.	1.1	54
3	IS ATTRACTION FATAL? THE EFFECTS OF HERBIVORE-INDUCED PLANT VOLATILES ON HERBIVORE PARASITISM. <i>Ecology</i> , 2002, 83, 3416-3425.	1.5	17
4	THE GENETIC ARCHITECTURE OF A COMPLEX ECOLOGICAL TRAIT: HOST PLANT USE IN THE SPECIALIST MOTH, <i>HELIOTHIS SUBFLEXA</i> . <i>Evolution; International Journal of Organic Evolution</i> , 2012, 66, 3336-3351.	1.1	15
5	Counties not countries: Variation in host specificity among populations of an aphid parasitoid. <i>Evolutionary Applications</i> , 2019, 12, 815-829.	1.5	15
6	The genetic architecture of ecological adaptation: intraspecific variation in host plant use by the lepidopteran crop pest <i>Chloridea virescens</i> . <i>Heredity</i> , 2018, 120, 234-250.	1.2	7
7	A whole genome gene content phylogenetic analysis of anopheline mosquitoes. <i>Molecular Phylogenetics and Evolution</i> , 2017, 107, 266-269.	1.2	6
8	De Novo characterization of transcriptomes from two North American <i>Papaipema</i> stem-borers (Lepidoptera: Noctuidae). <i>PLoS ONE</i> , 2018, 13, e0191061.	1.1	6
9	Transcriptome profiling with focus on potential key genes for wing development and evolution in <i>Megalopterus caeruleus</i> , the damselfly species with the world's largest wings. <i>PLoS ONE</i> , 2018, 13, e0189898.	1.1	4
10	Genome assembly of a Mesoamerican derived variety of lima bean: a foundational cultivar in the Mid-Atlantic USA. <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	0.8	4
11	Assemblies of the genomes of parasitic wasps using meta-assembly and scaffolding with genetic linkage. <i>G3: Genes, Genomes, Genetics</i> , 2021, , .	0.8	3
12	Genome content analysis yields new insights into the relationship between the human malaria parasite <i>Plasmodium falciparum</i> and its anopheline vectors. <i>BMC Genomics</i> , 2017, 18, 205.	1.2	2
13	Good Citizenship Made Easy: A Step-by-Step Guide to Submitting RNA-Seq Data to NCBI. <i>Current Protocols in Bioinformatics</i> , 2018, 64, e67.	25.8	1
14	The Agony and the Ecstasy: International Fieldwork from a Neophyte's Perspective. <i>American Entomologist</i> , 2018, 64, 88-93.	0.1	0