Mara K N Lawniczak

List of Publications by Year in descending order

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471509 677142 2,107 26 17 22 citations h-index g-index papers 36 36 36 3364 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A targeted amplicon sequencing panel to simultaneously identify mosquito species and <i>Plasmodium</i> presence across the entire <i>Anopheles</i> genus. Molecular Ecology Resources, 2022, 22, 28-44.	4.8	18
2	The Earth BioGenome Project 2020: Starting the clock. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119 , .	7.1	124
3	Standards recommendations for the Earth BioGenome Project. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	33
4	Why sequence all eukaryotes?. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	51
5	Sequence locally, think globally: The Darwin Tree of Life Project. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	120
6	Single-cell transcriptomics reveals expression profiles of Trypanosoma brucei sexual stages. PLoS Pathogens, 2022, 18, e1010346.	4.7	12
7	Tackling malaria transmission at a single cell level in an endemic setting in sub-Saharan Africa. Nature Communications, 2022, 13, 2679.	12.8	3
8	A Minimally Morphologically Destructive Approach for DNA Retrieval and Whole-Genome Shotgun Sequencing of Pinned Historic Dipteran Vector Species. Genome Biology and Evolution, 2021, 13, .	2.5	12
9	Using scRNA-seq to Identify Transcriptional Variation in the Malaria Parasite Ookinete Stage. Frontiers in Cellular and Infection Microbiology, 2021, 11, 604129.	3.9	8
10	A single-cell atlas of Plasmodium falciparum transmission through the mosquito. Nature Communications, 2021, 12, 3196.	12.8	54
11	Transmission of Artemisinin-Resistant Malaria Parasites to Mosquitoes under Antimalarial Drug Pressure. Antimicrobial Agents and Chemotherapy, 2020, 65, .	3.2	29
12	A microsporidian impairs Plasmodium falciparum transmission in Anopheles arabiensis mosquitoes. Nature Communications, 2020, 11, 2187.	12.8	62
13	Souporcell: robust clustering of single-cell RNA-seq data by genotype without reference genotypes. Nature Methods, 2020, 17, 615-620.	19.0	232
14	The Malaria Cell Atlas: Single parasite transcriptomes across the complete <i>Plasmodium</i> life cycle. Science, 2019, 365, .	12.6	198
15	Whole-genome sequencing reveals high complexity of copy number variation at insecticide resistance loci in malaria mosquitoes. Genome Research, 2019, 29, 1250-1261.	5.5	79
16	The male mosquito contribution towards malaria transmission: Mating influences the Anopheles female midgut transcriptome and increases female susceptibility to human malaria parasites. PLoS Pathogens, 2019, 15, e1008063.	4.7	25
17	A High-Quality De novo Genome Assembly from a Single Mosquito Using PacBio Sequencing. Genes, 2019, 10, 62.	2.4	121
18	Single-cell RNA-seq reveals hidden transcriptional variation in malaria parasites. ELife, 2018, 7, .	6.0	171

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19	Rapid evolution of female-biased genes among four species of <i>Anopheles</i> malaria mosquitoes. Genome Research, 2017, 27, 1536-1548.	5.5	60
20	Highly evolvable malaria vectors: The genomes of 16 <i>Anopheles</i> mosquitoes. Science, 2015, 347, 1258522.	12.6	492
21	Genetic Dissection of Anopheles gambiae Gut Epithelial Responses to Serratia marcescens. PLoS Pathogens, 2014, 10, e1003897.	4.7	87
22	Proteomic analysis of the Plasmodium male gamete reveals the key role for glycolysis in flagellar motility. Malaria Journal, 2014, 13, 315.	2.3	50
23	The genome sequence of the spiny starfish, Marthasterias glacialis (Linnaeus, 1758). Wellcome Open Research, 0, 6, 295.	1.8	3
24	The genome sequence of the blue-rayed limpet, Patella pellucida Linnaeus, 1758. Wellcome Open Research, 0, 7, 126.	1.8	2
25	Letâ \in ^M s talk about the (lady)birds and the bees: how insects can whisper a multitude of stories. ARPHA Conference Abstracts, 0, 5, .	0.0	0
26	Specimen and sample metadata standards for biodiversity genomics: a proposal from the Darwin Tree of Life project. Wellcome Open Research, 0, 7, 187.	1.8	11