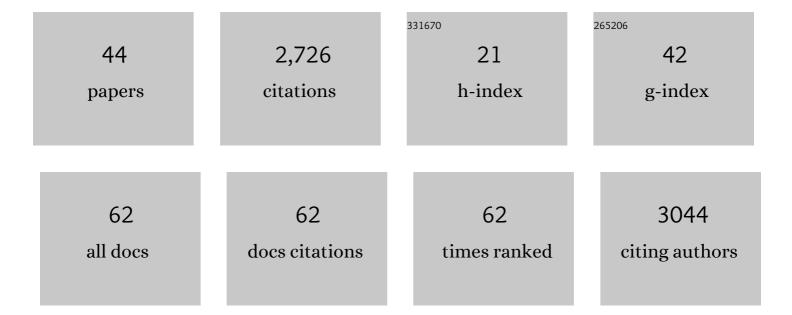
Thomas Walker

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4092945/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Association of Reduced Long-Lasting Insecticidal Net Efficacy and Pyrethroid Insecticide Resistance With Overexpression of <i>CYP6P4</i> , <i>CYP6P3,</i> and <i>CYP6Z1</i> in Populations of <i>Anopheles coluzzii</i> From Southeast Côte d'Ivoire. Journal of Infectious Diseases, 2022, 225, 1424-1434.	4.0	12
2	Wolbachia endosymbionts in two Anopheles species indicates independent acquisitions and lack of prophage elements. Microbial Genomics, 2022, 8, .	2.0	3
3	<i>Anopheles stephensi</i> Mosquitoes as Vectors of <i>Plasmodiumvivax</i> and <i>falciparum</i> , Horn of Africa, 2019. Emerging Infectious Diseases, 2021, 27, 603-607.	4.3	74
4	Evidence for natural hybridization and novel <i>Wolbachia</i> strain superinfections in the <i>Anopheles gambiae</i> complex from Guinea. Royal Society Open Science, 2021, 8, 202032.	2.4	11
5	Stable high-density and maternally inherited Wolbachia infections in Anopheles moucheti and Anopheles demeilloni mosquitoes. Current Biology, 2021, 31, 2310-2320.e5.	3.9	49
6	Overabundance of <i>Asaia</i> and <i>Serratia</i> Bacteria Is Associated with Deltamethrin Insecticide Susceptibility in <i>Anopheles coluzzii</i> from Agboville, Côte d'Ivoire. Microbiology Spectrum, 2021, 9, e0015721.	3.0	18
7	Investigating molecular mechanisms of insecticide resistance in the Eastern Democratic Republic of the Congo. Malaria Journal, 2021, 20, 464.	2.3	6
8	Habitat and Seasonality Affect Mosquito Community Composition in the West Region of Cameroon. Insects, 2020, 11, 312.	2.2	40
9	A Comparison of Adult Mosquito Trapping Methods to Assess Potential West Nile Virus Mosquito Vectors in Greece during the Onset of the 2018 Transmission Season. Insects, 2020, 11, 329.	2.2	5
10	An assessment of adult mosquito collection techniques for studying species abundance and diversity in Maferinyah, Guinea. Parasites and Vectors, 2020, 13, 150.	2.5	19
11	Evidence of extrinsic factors dominating intrinsic blood host preferences of major African malaria vectors. Scientific Reports, 2020, 10, 741.	3.3	13
12	Detection of a novel insect-specific flavivirus across ecologically diverse populations of Aedes aegypti on the Caribbean island of Saint Lucia. Wellcome Open Research, 2020, 5, 149.	1.8	0
13	Detection of Cell-Fusing Agent virus across ecologically diverse populations of Aedes aegypti on the Caribbean island of Saint Lucia. Wellcome Open Research, 2020, 5, 149.	1.8	4
14	Characterizing the molecular and metabolic mechanisms of insecticide resistance in Anopheles gambiae in Faranah, Guinea. Malaria Journal, 2019, 18, 244.	2.3	29
15	The relationship between insecticide resistance, mosquito age and malaria prevalence in Anopheles gambiae s.l. from Guinea. Scientific Reports, 2019, 9, 8846.	3.3	47
16	A community-level investigation following a yellow fever virus outbreak in South Omo Zone, South-West Ethiopia. PeerJ, 2019, 7, e6466.	2.0	12
17	Investigating the blood-host plasticity and dispersal of Anopheles coluzzii using a novel field-based methodology. Parasites and Vectors, 2019, 12, 143.	2.5	16
18	No evidence of Zika, dengue, or chikungunya virus infection in field-caught mosquitoes from the Recife Metropolitan Region, Brazil, 2015. Wellcome Open Research, 2019, 4, 93.	1.8	6

THOMAS WALKER

#	Article	IF	CITATIONS
19	Development of an urban molecular xenomonitoring system for lymphatic filariasis in the Recife Metropolitan Region, Brazil. PLoS Neglected Tropical Diseases, 2018, 12, e0006816.	3.0	10
20	Using the human blood index to investigate host biting plasticity: a systematic review and meta-regression of the three major African malaria vectors. Malaria Journal, 2018, 17, 479.	2.3	15
21	Diverse novel resident Wolbachia strains in Culicine mosquitoes from Madagascar. Scientific Reports, 2018, 8, 17456.	3.3	19
22	Novel Wolbachia strains in Anopheles malaria vectors from Sub-Saharan Africa. Wellcome Open Research, 2018, 3, 113.	1.8	34
23	Novel Wolbachia strains in Anopheles malaria vectors from Sub-Saharan Africa. Wellcome Open Research, 2018, 3, 113.	1.8	66
24	Establishment of a method for Lutzomyia longipalpis sand fly egg microinjection: The first step towards potential novel control strategies for leishmaniasis. Wellcome Open Research, 2018, 3, 55.	1.8	5
25	Biological Control of Mosquito Vectors: Past, Present, and Future. Insects, 2016, 7, 52.	2.2	255
26	Alternative vector control methods to manage the Zika virus outbreak: more haste, less speed – Authors' reply. The Lancet Global Health, 2016, 4, e365-e366.	6.3	3
27	Zika virus outbreak in the Americas: the need for novel mosquito control methods. The Lancet Global Health, 2016, 4, e148-e149.	6.3	144
28	Wolbachia Biocontrol Strategies for Arboviral Diseases and the Potential Influence of Resident Wolbachia Strains in Mosquitoes. Current Tropical Medicine Reports, 2016, 3, 20-25.	3.7	41
29	Establishment of a Wolbachia Superinfection in Aedes aegypti Mosquitoes as a Potential Approach for Future Resistance Management. PLoS Pathogens, 2016, 12, e1005434.	4.7	182
30	Comparison of Methods for Xenomonitoring in Vectors of Lymphatic Filariasis in Northeastern Tanzania. American Journal of Tropical Medicine and Hygiene, 2015, 93, 983-989.	1.4	11
31	The Potential Use of Wolbachia-Based Mosquito Biocontrol Strategies for Japanese Encephalitis. PLoS Neglected Tropical Diseases, 2015, 9, e0003576.	3.0	36
32	Limited Dengue Virus Replication in Field-Collected Aedes aegypti Mosquitoes Infected with Wolbachia. PLoS Neglected Tropical Diseases, 2014, 8, e2688.	3.0	288
33	Blood meal induced microRNA regulates development and immune associated genes in the Dengue mosquito vector, Aedes aegypti. Insect Biochemistry and Molecular Biology, 2013, 43, 146-152.	2.7	79
34	Genomic Evolution of the Pathogenic Wolbachia Strain, wMelPop. Genome Biology and Evolution, 2013, 5, 2189-2204.	2.5	96
35	Using bacteria to treat diseases. Expert Opinion on Biological Therapy, 2012, 12, 701-712.	3.1	11
36	Can Wolhachia be used to control malaria? Memorias Do Instituto Oswaldo Cruz, 2011, 106, 212-217	16	54

THOMAS WALKER

#	Article	IF	CITATIONS
37	<i>Wolbachia</i> and the biological control of mosquitoâ€borne disease. EMBO Reports, 2011, 12, 508-518.	4.5	349
38	Differentially expressed profiles in the larval testes of Wolbachia infected and uninfected Drosophila. BMC Genomics, 2011, 12, 595.	2.8	58
39	Wolbachia in the Culex pipiens Group Mosquitoes: Introgression and Superinfection. Journal of Heredity, 2009, 100, 192-196.	2.4	23
40	Horizontal gene transfer between Wolbachia and the mosquito Aedes aegypti. BMC Genomics, 2009, 10, 33.	2.8	142
41	Genome Evolution of Wolbachia Strain wPip from the Culex pipiens Group. Molecular Biology and Evolution, 2008, 25, 1877-1887.	8.9	210
42	Ankyrin repeat domain-encoding genes in the wPip strain of Wolbachia from the Culex pipiens group. BMC Biology, 2007, 5, 39.	3.8	60
43	Wolbachia variability and host effects on crossing type in Culex mosquitoes. Nature, 2005, 436, 257-260.	27.8	139
44	Establishment of a method for Lutzomyia longipalpis sand fly embryo microinjection: The first step	1.8	3

44 towards potential novel control strategies for leishmaniasis. Wellcome Open Research, 0, 3, 55.