

Thomas Walker

List of Publications by Year in descending order

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44
papers

2,726
citations

331670

21
h-index

265206

42
g-index

62
all docs

62
docs citations

62
times ranked

3044
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Wolbachia</i> and the biological control of mosquito-borne disease. <i>EMBO Reports</i> , 2011, 12, 508-518.	4.5	349
2	Limited Dengue Virus Replication in Field-Collected <i>Aedes aegypti</i> Mosquitoes Infected with <i>Wolbachia</i> . <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e2688.	3.0	288
3	Biological Control of Mosquito Vectors: Past, Present, and Future. <i>Insects</i> , 2016, 7, 52.	2.2	255
4	Genome Evolution of <i>Wolbachia</i> Strain wPip from the <i>Culex pipiens</i> Group. <i>Molecular Biology and Evolution</i> , 2008, 25, 1877-1887.	8.9	210
5	Establishment of a <i>Wolbachia</i> Superinfection in <i>Aedes aegypti</i> Mosquitoes as a Potential Approach for Future Resistance Management. <i>PLoS Pathogens</i> , 2016, 12, e1005434.	4.7	182
6	Zika virus outbreak in the Americas: the need for novel mosquito control methods. <i>The Lancet Global Health</i> , 2016, 4, e148-e149.	6.3	144
7	Horizontal gene transfer between <i>Wolbachia</i> and the mosquito <i>Aedes aegypti</i> . <i>BMC Genomics</i> , 2009, 10, 33.	2.8	142
8	<i>Wolbachia</i> variability and host effects on crossing type in <i>Culex</i> mosquitoes. <i>Nature</i> , 2005, 436, 257-260.	27.8	139
9	Genomic Evolution of the Pathogenic <i>Wolbachia</i> Strain, wMelPop. <i>Genome Biology and Evolution</i> , 2013, 5, 2189-2204.	2.5	96
10	Blood meal induced microRNA regulates development and immune associated genes in the Dengue mosquito vector, <i>Aedes aegypti</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2013, 43, 146-152.	2.7	79
11	<i>Anopheles stephensi</i> Mosquitoes as Vectors of <i>Plasmodium vivax</i> and <i>Plasmodium falciparum</i> , Horn of Africa, 2019. <i>Emerging Infectious Diseases</i> , 2021, 27, 603-607.	4.3	74
12	Novel <i>Wolbachia</i> strains in <i>Anopheles</i> malaria vectors from Sub-Saharan Africa. <i>Wellcome Open Research</i> , 2018, 3, 113.	1.8	66
13	Ankyrin repeat domain-encoding genes in the wPip strain of <i>Wolbachia</i> from the <i>Culex pipiens</i> group. <i>BMC Biology</i> , 2007, 5, 39.	3.8	60
14	Differentially expressed profiles in the larval testes of <i>Wolbachia</i> infected and uninfected <i>Drosophila</i> . <i>BMC Genomics</i> , 2011, 12, 595.	2.8	58
15	Can <i>Wolbachia</i> be used to control malaria?. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2011, 106, 212-217.	1.6	54
16	Stable high-density and maternally inherited <i>Wolbachia</i> infections in <i>Anopheles moucheti</i> and <i>Anopheles demeilloni</i> mosquitoes. <i>Current Biology</i> , 2021, 31, 2310-2320.e5.	3.9	49
17	The relationship between insecticide resistance, mosquito age and malaria prevalence in <i>Anopheles gambiae</i> s.l. from Guinea. <i>Scientific Reports</i> , 2019, 9, 8846.	3.3	47
18	<i>Wolbachia</i> Biocontrol Strategies for Arboviral Diseases and the Potential Influence of Resident <i>Wolbachia</i> Strains in Mosquitoes. <i>Current Tropical Medicine Reports</i> , 2016, 3, 20-25.	3.7	41

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19	Habitat and Seasonality Affect Mosquito Community Composition in the West Region of Cameroon. <i>Insects</i> , 2020, 11, 312.	2.2	40
20	The Potential Use of Wolbachia-Based Mosquito Biocontrol Strategies for Japanese Encephalitis. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003576.	3.0	36
21	Novel Wolbachia strains in Anopheles malaria vectors from Sub-Saharan Africa. <i>Wellcome Open Research</i> , 2018, 3, 113.	1.8	34
22	Characterizing the molecular and metabolic mechanisms of insecticide resistance in Anopheles gambiae in Faranah, Guinea. <i>Malaria Journal</i> , 2019, 18, 244.	2.3	29
23	Wolbachia in the Culex pipiens Group Mosquitoes: Introgression and Superinfection. <i>Journal of Heredity</i> , 2009, 100, 192-196.	2.4	23
24	Diverse novel resident Wolbachia strains in Culicine mosquitoes from Madagascar. <i>Scientific Reports</i> , 2018, 8, 17456.	3.3	19
25	An assessment of adult mosquito collection techniques for studying species abundance and diversity in Maferinyah, Guinea. <i>Parasites and Vectors</i> , 2020, 13, 150.	2.5	19
26	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. <i>Microbiology Spectrum</i> , 2021, 9, e0015721.	3.0	18
27	Investigating the blood-host plasticity and dispersal of Anopheles coluzzii using a novel field-based methodology. <i>Parasites and Vectors</i> , 2019, 12, 143.	2.5	16
28	Using the human blood index to investigate host biting plasticity: a systematic review and meta-regression of the three major African malaria vectors. <i>Malaria Journal</i> , 2018, 17, 479.	2.3	15
29	Evidence of extrinsic factors dominating intrinsic blood host preferences of major African malaria vectors. <i>Scientific Reports</i> , 2020, 10, 741.	3.3	13
30	A community-level investigation following a yellow fever virus outbreak in South Omo Zone, South-West Ethiopia. <i>PeerJ</i> , 2019, 7, e6466.	2.0	12
31	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. <i>Journal of Infectious Diseases</i> , 2022, 225, 1424-1434.	4.0	12
32	Using bacteria to treat diseases. <i>Expert Opinion on Biological Therapy</i> , 2012, 12, 701-712.	3.1	11
33	Comparison of Methods for Xenomonitoring in Vectors of Lymphatic Filariasis in Northeastern Tanzania. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015, 93, 983-989.	1.4	11
34	Evidence for natural hybridization and novel Wolbachia strain superinfections in the <i>Anopheles gambiae</i> complex from Guinea. <i>Royal Society Open Science</i> , 2021, 8, 202032.	2.4	11
35	Development of an urban molecular xenomonitoring system for lymphatic filariasis in the Recife Metropolitan Region, Brazil. <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006816.	3.0	10
36	No evidence of Zika, dengue, or chikungunya virus infection in field-caught mosquitoes from the Recife Metropolitan Region, Brazil, 2015. <i>Wellcome Open Research</i> , 2019, 4, 93.	1.8	6

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37	Investigating molecular mechanisms of insecticide resistance in the Eastern Democratic Republic of the Congo. <i>Malaria Journal</i> , 2021, 20, 464.	2.3	6
38	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. <i>Insects</i> , 2020, 11, 329.	2.2	5
39	Establishment of a method for <i>Lutzomyia longipalpis</i> sand fly egg microinjection: The first step towards potential novel control strategies for leishmaniasis. <i>Wellcome Open Research</i> , 2018, 3, 55.	1.8	5
40	Detection of Cell-Fusing Agent virus across ecologically diverse populations of <i>Aedes aegypti</i> on the Caribbean island of Saint Lucia. <i>Wellcome Open Research</i> , 2020, 5, 149.	1.8	4
41	Alternative vector control methods to manage the Zika virus outbreak: more haste, less speed – Authors' reply. <i>The Lancet Global Health</i> , 2016, 4, e365-e366.	6.3	3
42	Establishment of a method for <i>Lutzomyia longipalpis</i> sand fly embryo microinjection: The first step towards potential novel control strategies for leishmaniasis. <i>Wellcome Open Research</i> , 0, 3, 55.	1.8	3
43	<i>Wolbachia</i> endosymbionts in two <i>Anopheles</i> species indicates independent acquisitions and lack of prophage elements. <i>Microbial Genomics</i> , 2022, 8, .	2.0	3
44	Detection of a novel insect-specific flavivirus across ecologically diverse populations of <i>Aedes aegypti</i> on the Caribbean island of Saint Lucia. <i>Wellcome Open Research</i> , 2020, 5, 149.	1.8	0