Scott L O neill

List of Publications by Citations

Source: https://exaly.com/author-pdf/4373902/scott-l-oneill-publications-by-citations.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19,780 138 201 73 h-index g-index citations papers 206 6.71 22,767 7.4 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
201	A Wolbachia symbiont in Aedes aegypti limits infection with dengue, Chikungunya, and Plasmodium. <i>Cell</i> , 2009 , 139, 1268-78	56.2	1073
200	Successful establishment of Wolbachia in Aedes populations to suppress dengue transmission. <i>Nature</i> , 2011 , 476, 454-7	50.4	984
199	16S rRNA phylogenetic analysis of the bacterial endosymbionts associated with cytoplasmic incompatibility in insects. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1992 , 89, 2699-702	11.5	918
198	Phylogeny and PCR-based classification of Wolbachia strains using wsp gene sequences. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 1998 , 265, 509-15	4.4	880
197	The wMel Wolbachia strain blocks dengue and invades caged Aedes aegypti populations. <i>Nature</i> , 2011 , 476, 450-3	50.4	841
196	Wolbachia and virus protection in insects. <i>Science</i> , 2008 , 322, 702	33.3	799
195	Stable introduction of a life-shortening Wolbachia infection into the mosquito Aedes aegypti. <i>Science</i> , 2009 , 323, 141-4	33.3	656
194	Phylogenomics of the reproductive parasite Wolbachia pipientis wMel: a streamlined genome overrun by mobile genetic elements. <i>PLoS Biology</i> , 2004 , 2, E69	9.7	613
193	Cloning and characterization of a gene encoding the major surface protein of the bacterial endosymbiont Wolbachia pipientis. <i>Journal of Bacteriology</i> , 1998 , 180, 2373-8	3.5	472
192	Bidirectional incompatibility between conspecific populations of Drosophila simulans. <i>Nature</i> , 1990 , 348, 178-80	50.4	340
191	Wolbachia infections are distributed throughout insect somatic and germ line tissues. <i>Insect Biochemistry and Molecular Biology</i> , 1999 , 29, 153-60	4.5	299
190	Interspecific and intraspecific horizontal transfer of Wolbachia in Drosophila. <i>Science</i> , 1993 , 260, 1796-9	33.3	294
189	Wolbachia and the biological control of mosquito-borne disease. <i>EMBO Reports</i> , 2011 , 12, 508-18	6.5	269
188	Beyond insecticides: new thinking on an ancient problem. <i>Nature Reviews Microbiology</i> , 2013 , 11, 181-9	322.2	257
187	Variation in antiviral protection mediated by different Wolbachia strains in Drosophila simulans. <i>PLoS Pathogens</i> , 2009 , 5, e1000656	7.6	257
186	Evidence for metabolic provisioning by a common invertebrate endosymbiont, Wolbachia pipientis, during periods of nutritional stress. <i>PLoS Pathogens</i> , 2009 , 5, e1000368	7.6	254
185	Impact of Wolbachia on infection with chikungunya and yellow fever viruses in the mosquito vector Aedes aegypti. <i>PLoS Neglected Tropical Diseases</i> , 2012 , 6, e1892	4.8	247

(2010-2002)

184	Wolbachia density and virulence attenuation after transfer into a novel host. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002 , 99, 2918-23	11.5	237	
183	Limited dengue virus replication in field-collected Aedes aegypti mosquitoes infected with Wolbachia. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e2688	4.8	229	
182	The relative importance of innate immune priming in Wolbachia-mediated dengue interference. <i>PLoS Pathogens</i> , 2012 , 8, e1002548	7.6	214	
181	Stability of the wMel Wolbachia Infection following invasion into Aedes aegypti populations. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e3115	4.8	204	
180	Wolbachia superinfections and the expression of cytoplasmic incompatibility. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 1995 , 261, 325-30	4.4	192	
179	Evidence for a global Wolbachia replacement in Drosophila melanogaster. <i>Current Biology</i> , 2005 , 15, 1428-33	6.3	181	
178	Genome evolution of Wolbachia strain wPip from the Culex pipiens group. <i>Molecular Biology and Evolution</i> , 2008 , 25, 1877-87	8.3	179	
177	Dietary cholesterol modulates pathogen blocking by Wolbachia. <i>PLoS Pathogens</i> , 2013 , 9, e1003459	7.6	177	
176	Wolbachia uses host microRNAs to manipulate host gene expression and facilitate colonization of the dengue vector Aedes aegypti. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 9250-5	11.5	176	
175	Modeling the impact on virus transmission of Wolbachia-mediated blocking of dengue virus infection of Aedes aegypti. <i>Science Translational Medicine</i> , 2015 , 7, 279ra37	17.5	165	
174	Antiviral protection and the importance of Wolbachia density and tissue tropism in Drosophila simulans. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 6922-9	4.8	156	
173	Replacement of the natural Wolbachia symbiont of Drosophila simulans with a mosquito counterpart. <i>Nature</i> , 1994 , 367, 453-5	50.4	156	
172	Local introduction and heterogeneous spatial spread of dengue-suppressing Wolbachia through an urban population of Aedes aegypti. <i>PLoS Biology</i> , 2017 , 15, e2001894	9.7	155	
171	Wolbachia uses a host microRNA to regulate transcripts of a methyltransferase, contributing to dengue virus inhibition in Aedes aegypti. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 10276-81	11.5	152	
170	Controlling vector-borne diseases by releasing modified mosquitoes. <i>Nature Reviews Microbiology</i> , 2018 , 16, 508-518	22.2	150	
169	Scaled deployment of Wolbachia to protect the community from dengue and other Aedes transmitted arboviruses. <i>Gates Open Research</i> , 2018 , 2, 36	2.4	147	
168	Distribution and diversity of Wolbachia infections in Southeast Asian mosquitoes (Diptera: Culicidae). <i>Journal of Medical Entomology</i> , 2000 , 37, 340-5	2.2	146	
167	Wolbachia-mediated resistance to dengue virus infection and death at the cellular level. <i>PLoS ONE</i> , 2010 , 5, e13398	3.7	142	

166	Rescuing Wolbachia have been overlooked. <i>Nature</i> , 1998 , 391, 852-3	50.4	142
165	Establishment of a Wolbachia Superinfection in Aedes aegypti Mosquitoes as a Potential Approach for Future Resistance Management. <i>PLoS Pathogens</i> , 2016 , 12, e1005434	7.6	142
164	Taxonomic status of the intracellular bacterium Wolbachia pipientis. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007 , 57, 654-657	2.2	140
163	In vitro cultivation of Wolbachia pipientis in an Aedes albopictus cell line. <i>Insect Molecular Biology</i> , 1997 , 6, 33-9	3.4	139
162	Wolbachia infections and the expression of cytoplasmic incompatibility in Drosophila sechellia and D. mauritiana. <i>Genetics</i> , 1995 , 140, 1307-17	4	138
161	Dynamics of the "popcorn" Wolbachia infection in outbred Aedes aegypti informs prospects for mosquito vector control. <i>Genetics</i> , 2011 , 187, 583-95	4	133
160	Tissue distribution and prevalence of Wolbachia infections in tsetse flies, Glossina spp. <i>Medical and Veterinary Entomology</i> , 2000 , 14, 44-50	2.4	132
159	Field evaluation of the establishment potential of wMelPop Wolbachia in Australia and Vietnam for dengue control. <i>Parasites and Vectors</i> , 2015 , 8, 563	4	128
158	Wolbachia infection reduces blood-feeding success in the dengue fever mosquito, Aedes aegypti. <i>PLoS Neglected Tropical Diseases</i> , 2009 , 3, e516	4.8	128
157	Modification of arthropod vector competence via symbiotic bacteria. <i>Parasitology Today</i> , 1993 , 9, 179-	83	126
157 156	Modification of arthropod vector competence via symbiotic bacteria. <i>Parasitology Today</i> , 1993 , 9, 179- Wolbachia neither induces nor suppresses transcripts encoding antimicrobial peptides. <i>Insect Molecular Biology</i> , 2000 , 9, 635-9	3.4	126
	Wolbachia neither induces nor suppresses transcripts encoding antimicrobial peptides. <i>Insect</i>		
156	Wolbachia neither induces nor suppresses transcripts encoding antimicrobial peptides. <i>Insect Molecular Biology</i> , 2000 , 9, 635-9 Distribution, expression, and motif variability of ankyrin domain genes in Wolbachia pipientis.	3.4	120
156 155	Wolbachia neither induces nor suppresses transcripts encoding antimicrobial peptides. <i>Insect Molecular Biology</i> , 2000 , 9, 635-9 Distribution, expression, and motif variability of ankyrin domain genes in Wolbachia pipientis. <i>Journal of Bacteriology</i> , 2005 , 187, 5136-45 Scaled deployment of to protect the community from dengue and other 'transmitted arboviruses.	3.4	120
156 155 154	Wolbachia neither induces nor suppresses transcripts encoding antimicrobial peptides. <i>Insect Molecular Biology</i> , 2000 , 9, 635-9 Distribution, expression, and motif variability of ankyrin domain genes in Wolbachia pipientis. <i>Journal of Bacteriology</i> , 2005 , 187, 5136-45 Scaled deployment of to protect the community from dengue and other transmitted arboviruses. <i>Gates Open Research</i> , 2018 , 2, 36 Host adaptation of a Wolbachia strain after long-term serial passage in mosquito cell lines. <i>Applied</i>	3·4 3·5 2·4	120 114 114
156 155 154 153	Wolbachia neither induces nor suppresses transcripts encoding antimicrobial peptides. <i>Insect Molecular Biology</i> , 2000 , 9, 635-9 Distribution, expression, and motif variability of ankyrin domain genes in Wolbachia pipientis. <i>Journal of Bacteriology</i> , 2005 , 187, 5136-45 Scaled deployment of to protect the community from dengue and other transmitted arboviruses. <i>Gates Open Research</i> , 2018 , 2, 36 Host adaptation of a Wolbachia strain after long-term serial passage in mosquito cell lines. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 6963-9 A virulent Wolbachia infection decreases the viability of the dengue vector Aedes aegypti during	3·4 3·5 2·4 4.8	120 114 114 112
156 155 154 153	Wolbachia neither induces nor suppresses transcripts encoding antimicrobial peptides. <i>Insect Molecular Biology</i> , 2000 , 9, 635-9 Distribution, expression, and motif variability of ankyrin domain genes in Wolbachia pipientis. <i>Journal of Bacteriology</i> , 2005 , 187, 5136-45 Scaled deployment of to protect the community from dengue and other 'transmitted arboviruses. <i>Gates Open Research</i> , 2018 , 2, 36 Host adaptation of a Wolbachia strain after long-term serial passage in mosquito cell lines. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 6963-9 A virulent Wolbachia infection decreases the viability of the dengue vector Aedes aegypti during periods of embryonic quiescence. <i>PLoS Neglected Tropical Diseases</i> , 2010 , 4, e748 Phylogenetically distant symbiotic microorganisms reside in Glossina midgut and ovary tissues.	3.4 3.5 2.4 4.8	120 114 114 112 110

(2009-1995)

148	Wolbachia pipientis: bacterial density and unidirectional cytoplasmic incompatibility between infected populations of Aedes albopictus. <i>Experimental Parasitology</i> , 1995 , 81, 284-91	2.1	101
147	Wolbachia Reduces the Transmission Potential of Dengue-Infected Aedes aegypti. <i>PLoS Neglected Tropical Diseases</i> , 2015 , 9, e0003894	4.8	94
146	Competition for amino acids between Wolbachia and the mosquito host, Aedes aegypti. <i>Microbial Ecology</i> , 2014 , 67, 205-18	4.4	91
145	Establishment of Mel in mosquitoes and reduction of local dengue transmission in Cairns and surrounding locations in northern Queensland, Australia. <i>Gates Open Research</i> , 2019 , 3, 1547	2.4	88
144	Wolbachia-associated bacterial protection in the mosquito Aedes aegypti. <i>PLoS Neglected Tropical Diseases</i> , 2013 , 7, e2362	4.8	87
143	Genetic transformation and phylogeny of bacterial symbionts from tsetse. <i>Insect Molecular Biology</i> , 1993 , 1, 123-31	3.4	86
142	An ancient horizontal gene transfer between mosquito and the endosymbiotic bacterium Wolbachia pipientis. <i>Molecular Biology and Evolution</i> , 2009 , 26, 367-74	8.3	84
141	Modifying insect population age structure to control vector-borne disease. <i>Advances in Experimental Medicine and Biology</i> , 2008 , 627, 126-40	3.6	83
140	Male development time influences the strength of Wolbachia-induced cytoplasmic incompatibility expression in Drosophila melanogaster. <i>Genetics</i> , 2007 , 177, 801-8	4	81
139	Wolbachia pipientis: intracellular infection and pathogenesis in Drosophila. <i>Current Opinion in Microbiology</i> , 2004 , 7, 67-70	7.9	81
138	Wolbachia-mediated sperm modification is dependent on the host genotype in Drosophila. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2001 , 268, 2565-70	4.4	81
137	The use of transcriptional profiles to predict adult mosquito age under field conditions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 18060-5	11.5	80
136	Guidance for contained field trials of vector mosquitoes engineered to contain a gene drive system: recommendations of a scientific working group. <i>Vector-Borne and Zoonotic Diseases</i> , 2008 , 8, 127-66	2.4	79
135	Mutual exclusion of Asaia and Wolbachia in the reproductive organs of mosquito vectors. <i>Parasites and Vectors</i> , 2015 , 8, 278	4	77
134	Genomic evolution of the pathogenic Wolbachia strain, wMelPop. <i>Genome Biology and Evolution</i> , 2013 , 5, 2189-204	3.9	77
133	Human probing behavior of Aedes aegypti when infected with a life-shortening strain of Wolbachia. <i>PLoS Neglected Tropical Diseases</i> , 2009 , 3, e568	4.8	77
132	Matching the genetics of released and local Aedes aegypti populations is critical to assure Wolbachia invasion. <i>PLoS Neglected Tropical Diseases</i> , 2019 , 13, e0007023	4.8	77
131	Increased locomotor activity and metabolism of Aedes aegypti infected with a life-shortening strain of Wolbachia pipientis. <i>Journal of Experimental Biology</i> , 2009 , 212, 1436-41	3	76

130	Field- and clinically derived estimates of -mediated blocking of dengue virus transmission potential in mosquitoes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, 361-366	11.5	75
129	Establishment of wMel Wolbachia in Aedes aegypti mosquitoes and reduction of local dengue transmission in Cairns and surrounding locations in northern Queensland, Australia. <i>Gates Open Research</i> , 2019 , 3, 1547	2.4	75
128	Characterization of Wolbachia host cell range via the in vitro establishment of infections. <i>Applied and Environmental Microbiology</i> , 2002 , 68, 656-60	4.8	73
127	A stable triple Wolbachia infection in Drosophila with nearly additive incompatibility effects. <i>Heredity</i> , 1999 , 82 (Pt 6), 620-7	3.6	72
126	Determination of Wolbachia genome size by pulsed-field gel electrophoresis. <i>Journal of Bacteriology</i> , 2001 , 183, 2219-25	3.5	71
125	wsp gene sequences from the Wolbachia of filarial nematodes. Current Microbiology, 2000 , 41, 96-100	2.4	71
124	Blood meal induced microRNA regulates development and immune associated genes in the Dengue mosquito vector, Aedes aegypti. <i>Insect Biochemistry and Molecular Biology</i> , 2013 , 43, 146-52	4.5	67
123	Novel Wolbachia-transinfected Aedes aegypti mosquitoes possess diverse fitness and vector competence phenotypes. <i>PLoS Pathogens</i> , 2017 , 13, e1006751	7.6	66
122	The Use of Wolbachia by the World Mosquito Program to Interrupt Transmission of Aedes aegypti Transmitted Viruses. <i>Advances in Experimental Medicine and Biology</i> , 2018 , 1062, 355-360	3.6	66
121	Prospects for control of African trypanosomiasis by tsetse vector manipulation. <i>Trends in Parasitology</i> , 2001 , 17, 29-35	6.4	65
12 0	Wolbachia infections of tephritid fruit flies: molecular evidence for five distinct strains in a single host species. <i>Current Microbiology</i> , 2002 , 45, 255-60	2.4	62
119	Francisella-like endosymbionts of ticks. <i>Journal of Invertebrate Pathology</i> , 2000 , 76, 301-3	2.6	62
118	Distribution and Diversity of Wolbachia Infections in Southeast Asian Mosquitoes (Diptera: Culicidae). <i>Journal of Medical Entomology</i> , 2000 , 37, 340-345	2.2	61
117	The toll and Imd pathways are not required for wolbachia-mediated dengue virus interference. <i>Journal of Virology</i> , 2013 , 87, 11945-9	6.6	60
116	Efficacy of Wolbachia-Infected Mosquito Deployments for the Control of Dengue. <i>New England Journal of Medicine</i> , 2021 , 384, 2177-2186	59.2	59
115	Field prevalence of Wolbachia in the mosquito vector Aedes albopictus. <i>American Journal of Tropical Medicine and Hygiene</i> , 2002 , 66, 108-11	3.2	58
114	Host age effect and expression of cytoplasmic incompatibility in field populations of Wolbachia-superinfected Aedes albopictus. <i>Heredity</i> , 2002 , 88, 270-4	3.6	56
113	Wolbachia small noncoding RNAs and their role in cross-kingdom communications. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 18721-6	11.5	55

(2011-2010)

112	Assessing key safety concerns of a Wolbachia-based strategy to control dengue transmission by Aedes mosquitoes. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2010 , 105, 957-64	2.6	54
111	Maternal transmission efficiency of Wolbachia superinfections in Aedes albopictus populations in Thailand. <i>American Journal of Tropical Medicine and Hygiene</i> , 2002 , 66, 103-7	3.2	54
110	Wolbachia infections of phlebotomine sand flies (Diptera: Psychodidae). <i>Journal of Medical Entomology</i> , 2001 , 38, 237-41	2.2	53
109	Assessing the epidemiological effect of wolbachia for dengue control. <i>Lancet Infectious Diseases, The</i> , 2015 , 15, 862-6	25.5	52
108	Insect densoviruses may be widespread in mosquito cell lines. <i>Journal of General Virology</i> , 1995 , 76 (Pt 8), 2067-74	4.9	51
107	Draft genome sequence of the male-killing Wolbachia strain wBol1 reveals recent horizontal gene transfers from diverse sources. <i>BMC Genomics</i> , 2013 , 14, 20	4.5	49
106	High anti-viral protection without immune upregulation after interspecies Wolbachia transfer. <i>PLoS ONE</i> , 2014 , 9, e99025	3.7	46
105	Infection with a Virulent Strain of Wolbachia Disrupts Genome Wide-Patterns of Cytosine Methylation in the Mosquito Aedes aegypti. <i>PLoS ONE</i> , 2013 , 8, e66482	3.7	46
104	A secure semi-field system for the study of Aedes aegypti. <i>PLoS Neglected Tropical Diseases</i> , 2011 , 5, e988	4.8	46
103	Reduced dengue incidence following deployments of -infected in Yogyakarta, Indonesia: a quasi-experimental trial using controlled interrupted time series analysis. <i>Gates Open Research</i> , 2020 , 4, 50	2.4	46
102	Wolbachia-induced aae-miR-12 miRNA negatively regulates the expression of MCT1 and MCM6 genes in Wolbachia-infected mosquito cell line. <i>PLoS ONE</i> , 2012 , 7, e50049	3.7	46
101	The AWED trial (Applying Wolbachia to Eliminate Dengue) to assess the efficacy of Wolbachia-infected mosquito deployments to reduce dengue incidence in Yogyakarta, Indonesia: study protocol for a cluster randomised controlled trial. <i>Trials</i> , 2018 , 19, 302	2.8	42
100	A Wolbachia symbiont in Aedes aegypti disrupts mosquito egg development to a greater extent when mosquitoes feed on nonhuman versus human blood. <i>Journal of Medical Entomology</i> , 2011 , 48, 76-	·8 ² 4 ²	41
99	Stable establishment of wMel Wolbachia in Aedes aegypti populations in Yogyakarta, Indonesia. <i>PLoS Neglected Tropical Diseases</i> , 2020 , 14, e0008157	4.8	41
98	Comparison of Stable and Transient Wolbachia Infection Models in Aedes aegypti to Block Dengue and West Nile Viruses. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005275	4.8	39
97	Wolbachia infection alters the relative abundance of resident bacteria in adult Aedes aegypti mosquitoes, but not larvae. <i>Molecular Ecology</i> , 2018 , 27, 297-309	5.7	38
96	Wolbachia-host interactions: connecting phenotype to genotype. <i>Current Opinion in Microbiology</i> , 2007 , 10, 221-4	7.9	37
95	Functional test of the influence of Wolbachia genes on cytoplasmic incompatibility expression in Drosophila melanogaster. <i>Insect Molecular Biology</i> , 2011 , 20, 75-85	3.4	36

94	Tandem repeat markers as novel diagnostic tools for high resolution fingerprinting of Wolbachia. <i>BMC Microbiology</i> , 2012 , 12 Suppl 1, S12	4.5	35
93	Structural and functional characterization of the oxidoreductase alpha-DsbA1 from Wolbachia pipientis. <i>Antioxidants and Redox Signaling</i> , 2009 , 11, 1485-500	8.4	35
92	Wolbachia Infection in the Coffee Berry Borer (Coleoptera: Scolytidae). <i>Annals of the Entomological Society of America</i> , 2002 , 95, 374-378	2	35
91	A mosquito densovirus infecting Aedes aegypti and Aedes albopictus from Thailand. <i>American Journal of Tropical Medicine and Hygiene</i> , 1999 , 61, 612-7	3.2	35
90	Wolbachia interferes with the intracellular distribution of Argonaute 1 in the dengue vector Aedes aegypti by manipulating the host microRNAs. <i>RNA Biology</i> , 2013 , 10, 1868-75	4.8	34
89	Variable infection frequency and high diversity of multiple strains of Wolbachia pipientis in Perkinsiella Planthoppers. <i>Applied and Environmental Microbiology</i> , 2011 , 77, 2165-8	4.8	34
88	wMel limits zika and chikungunya virus infection in a Singapore Wolbachia-introgressed Ae. aegypti strain, wMel-Sg. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005496	4.8	34
87	Infection with the wMel and wMelPop strains of Wolbachia leads to higher levels of melanization in the hemolymph of Drosophila melanogaster, Drosophila simulans and Aedes aegypti. Developmental and Comparative Immunology, 2011, 35, 360-5	3.2	32
86	The small interfering RNA pathway is not essential for Wolbachia-mediated antiviral protection in Drosophila melanogaster. <i>Applied and Environmental Microbiology</i> , 2012 , 78, 6773-6	4.8	31
85	Predicting the age of mosquitoes using transcriptional profiles. <i>Nature Protocols</i> , 2007 , 2, 2796-806	18.8	31
84	Wolbachia infection and expression of cytoplasmic incompatibility in Armigeres subalbatus (Diptera: Culicidae). <i>Journal of Medical Entomology</i> , 2000 , 37, 53-7	2.2	31
83	Evolution of Wolbachia pipientis transmission dynamics in insects. <i>Trends in Microbiology</i> , 1999 , 7, 297-	302 .4	30
82	Wolbachia-mediated virus blocking in mosquito cells is dependent on XRN1-mediated viral RNA degradation and influenced by viral replication rate. <i>PLoS Pathogens</i> , 2018 , 14, e1006879	7.6	29
81	Why do we need alternative tools to control mosquito-borne diseases in Latin America?. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2012 , 107, 828-9	2.6	29
80	Development of a physical and genetic map of the virulent Wolbachia strain wMelPop. <i>Journal of Bacteriology</i> , 2003 , 185, 7077-84	3.5	28
79	Wolbachia genomes: insights into an intracellular lifestyle. <i>Current Biology</i> , 2005 , 15, R507-9	6.3	27
78	Crossing type variability associated with cytoplasmic incompatibility in Australian populations of the mosquito Culex quinquefasciatus Say. <i>Medical and Veterinary Entomology</i> , 1992 , 6, 209-16	2.4	27
77	Cytoplasmic symbionts in Tribolium confusum. <i>Journal of Invertebrate Pathology</i> , 1989 , 53, 132-134	2.6	26

76	Influence of the virus LbFV and of Wolbachia in a host-parasitoid interaction. <i>PLoS ONE</i> , 2012 , 7, e3508	1 3.7	24
75	Molecular phylogeny of Wolbachia endosymbionts in Southeast Asian mosquitoes (Diptera: Culicidae) based on wsp gene sequences. <i>Journal of Medical Entomology</i> , 2003 , 40, 1-5	2.2	24
74	Beyond the Q ack yard Q Lay knowledge about Aedes aegypti in northern Australia and its implications for policy and practice. <i>Acta Tropica</i> , 2010 , 116, 74-80	3.2	23
73	Absence of the symbiont Candidatus Midichloria mitochondrii in the mitochondria of the tick Ixodes holocyclus. <i>FEMS Microbiology Letters</i> , 2009 , 299, 241-7	2.9	23
72	Comparative susceptibility of mosquito populations in North Queensland, Australia to oral infection with dengue virus. <i>American Journal of Tropical Medicine and Hygiene</i> , 2014 , 90, 422-30	3.2	22
71	Transinfected Wolbachia have minimal effects on male reproductive success in Aedes aegypti. <i>Parasites and Vectors</i> , 2013 , 6, 36	4	22
70	The wMelPop strain of Wolbachia interferes with dopamine levels in Aedes aegypti. <i>Parasites and Vectors</i> , 2011 , 4, 28	4	22
69	Multiple Wolbachia strains provide comparative levels of protection against dengue virus infection in Aedes aegypti. <i>PLoS Pathogens</i> , 2020 , 16, e1008433	7.6	21
68	Rapid spread of male-killing Wolbachia in the butterfly Hypolimnas bolina. <i>Journal of Evolutionary Biology</i> , 2010 , 23, 231-5	2.3	21
67	Detecting wMel Wolbachia in field-collected Aedes aegypti mosquitoes using loop-mediated isothermal amplification (LAMP). <i>Parasites and Vectors</i> , 2019 , 12, 404	4	20
66	Impacts of Wolbachia infection on predator prey relationships: evaluating survival and horizontal transfer between wMelPop infected Aedes aegypti and its predators. <i>Journal of Medical Entomology</i> , 2012 , 49, 624-30	2.2	20
65	Scaled deployment of Wolbachia to protect the community from Aedes transmitted arboviruses. Gates Open Research, 2, 36	2.4	20
64	A simple protocol to obtain highly pure Wolbachia endosymbiont DNA for genome sequencing. Journal of Microbiological Methods, 2011 , 84, 134-6	2.8	19
63	Wolbachia replication and host cell division in Aedes albopictus. <i>Current Microbiology</i> , 2004 , 49, 10-2	2.4	19
62	Epidemiological, Serological, and Virological Features of Dengue in Nha Trang City, Vietnam. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018 , 98, 402-409	3.2	19
61	Screening of Wolbachia Endosymbiont Infection in Aedes aegypti Mosquitoes Using Attenuated Total Reflection Mid-Infrared Spectroscopy. <i>Analytical Chemistry</i> , 2017 , 89, 5285-5293	7.8	18
60	Evolutionary dynamics of insect symbiont associations. <i>Trends in Ecology and Evolution</i> , 2007 , 22, 625-7	10.9	18
59	Field validation of a transcriptional assay for the prediction of age of uncaged Aedes aegypti mosquitoes in Northern Australia. <i>PLoS Neglected Tropical Diseases</i> , 2010 , 4, e608	4.8	18

58	"Endomicrobia" and other bacteria associated with the hindgut of Dermolepida albohirtum larvae. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 762-7	4.8	17
57	Assessment of gut bacteria for a paratransgenic approach to control Dermolepida albohirtum larvae. <i>Applied and Environmental Microbiology</i> , 2008 , 74, 4036-43	4.8	17
56	Wolbachia pipientis: symbiont or parasite?. <i>Parasitology Today</i> , 1995 , 11, 168-9		17
55	Evidence of a spotted fever-like rickettsia and a potential new vector from northeastern Australia. <i>Journal of Medical Entomology</i> , 2005 , 42, 918-21	2.2	16
54	SOCIAL BEHAVIOUR AND ITS RELATIONSHIP TO FIELD DISTRIBUTION IN PANESTHIA CRIBRATA SAUSSURE (BLATTODEA: BLABERIDAE). <i>Australian Journal of Entomology</i> , 1987 , 26, 313-321		16
53	Effectiveness of Wolbachia-infected mosquito deployments in reducing the incidence of dengue and other Aedes-borne diseases in Niter[] Brazil: A quasi-experimental study. <i>PLoS Neglected Tropical Diseases</i> , 2021 , 15, e0009556	4.8	16
52	A portable approach for the surveillance of dengue virus-infected mosquitoes. <i>Journal of Virological Methods</i> , 2012 , 183, 90-3	2.6	15
51	Improved accuracy of the transcriptional profiling method of age grading in Aedes aegypti mosquitoes under laboratory and semi-field cage conditions and in the presence of Wolbachia infection. <i>Insect Molecular Biology</i> , 2011 , 20, 215-24	3.4	15
50	Control of Vector-Borne Disease by Genetic Manipulation of Insect Populations: Technological Requirements and Research Priorities. <i>Australian Journal of Entomology</i> , 1997 , 36, 309-317		15
49	Analysis of Wolbachia protein synthesis in Drosophila in vivo. <i>Insect Molecular Biology</i> , 1998 , 7, 101-5	3.4	15
48	Intensity of Mutualism Breakdown Is Determined by Temperature Not Amplification of Wolbachia Genes. <i>PLoS Pathogens</i> , 2016 , 12, e1005888	7.6	15
47	Novel phenotype of Wolbachia strain wPip in Aedes aegypti challenges assumptions on mechanisms of Wolbachia-mediated dengue virus inhibition. <i>PLoS Pathogens</i> , 2020 , 16, e1008410	7.6	15
46	Wolbachia introduction into Lutzomyia longipalpis (Diptera: Psychodidae) cell lines and its effects on immune-related gene expression and interaction with Leishmania infantum. <i>Parasites and Vectors</i> , 2019 , 12, 33	4	15
45	Identification of yeast associated with the planthopper, Perkinsiella saccharicida: potential applications for Fiji leaf gall control. <i>Current Microbiology</i> , 2011 , 63, 392-401	2.4	14
44	Update to the AWED (Applying Wolbachia to Eliminate Dengue) trial study protocol: a cluster randomised controlled trial in Yogyakarta, Indonesia. <i>Trials</i> , 2020 , 21, 429	2.8	13
43	A Native Wolbachia Endosymbiont Does Not Limit Dengue Virus Infection in the Mosquito Aedes notoscriptus (Diptera: Culicidae). <i>Journal of Medical Entomology</i> , 2016 , 53, 401-8	2.2	13
42	Baseline Characterization of Dengue Epidemiology in Yogyakarta City, Indonesia, before a Randomized Controlled Trial of for Arboviral Disease Control. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018 , 99, 1299-1307	3.2	13
41	A highly stable blood meal alternative for rearing Aedes and Anopheles mosquitoes. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0006142	4.8	12

(2020-2018)

40	Cluster-Randomized Test-Negative Design Trials: A Novel and Efficient Method to Assess the Efficacy of Community-Level Dengue Interventions. <i>American Journal of Epidemiology</i> , 2018 , 187, 2021	1-2028	12
39	A Rapid Single-Step Multiplex Method for Discriminating Between Trichogramma (Hymenoptera: Trichogrammatidae) Species in Australia. <i>Journal of Economic Entomology</i> , 2006 , 99, 2142-2145	2.2	12
38	Wolbachia mosquito control: Tested. <i>Science</i> , 2016 , 352, 526	33.3	11
37	Spatial and Temporal Variation in Aedes aegypti and Aedes albopictus (Diptera: Culicidae) Numbers in the Yogyakarta Area of Java, Indonesia, With Implications for Wolbachia Releases. <i>Journal of Medical Entomology</i> , 2016 , 53, 188-98	2.2	10
36	Differential suppression of persistent insect specific viruses in trans-infected wMel and wMelPop-CLA Aedes-derived mosquito lines. <i>Virology</i> , 2019 , 527, 141-145	3.6	10
35	New names for old strains? Wolbachia wSim is actually wRi. <i>Genome Biology</i> , 2005 , 6, 401; author reply 401	18.3	8
34	Cytoplasmic incompatibility in Drosophila populations: influence of assortative mating on symbiont distribution. <i>Journal of Invertebrate Pathology</i> , 1991 , 58, 436-43	2.6	8
33	How to engage communities on a large scale? Lessons from World Mosquito Program in Rio de Janeiro, Brazil. <i>Gates Open Research</i> , 2020 , 4, 109	2.4	8
32	Cloning, expression, purification and characterization of a DsbA-like protein from Wolbachia pipientis. <i>Protein Expression and Purification</i> , 2008 , 59, 266-73	2	7
31	THE DENGUE STOPPER. Scientific American, 2015 , 312, 72-7	0.5	6
30	Investigation of environmental influences on a transcriptional assay for the prediction of age of Aedes aegypti (Diptera: Culicidae) mosquitoes. <i>Journal of Medical Entomology</i> , 2010 , 47, 1044-52	2.2	6
29		2.2 3.6	6
	Aedes aegypti (Diptera: Culicidae) mosquitoes. <i>Journal of Medical Entomology</i> , 2010 , 47, 1044-52 The impact of large-scale deployment of mosquitoes on arboviral disease incidence in Rio de Janeiro and Niter[] Brazil: study protocol for a controlled interrupted time series analysis using		
29	Aedes aegypti (Diptera: Culicidae) mosquitoes. <i>Journal of Medical Entomology</i> , 2010 , 47, 1044-52 The impact of large-scale deployment of mosquitoes on arboviral disease incidence in Rio de Janeiro and Niter[] Brazil: study protocol for a controlled interrupted time series analysis using routine disease surveillance data. <i>F1000Research</i> , 2019 , 8, 1328 Large-Scale Deployment and Establishment of Into the Population in Rio de Janeiro, Brazil.	3.6	6
29	Aedes aegypti (Diptera: Culicidae) mosquitoes. <i>Journal of Medical Entomology</i> , 2010 , 47, 1044-52 The impact of large-scale deployment of mosquitoes on arboviral disease incidence in Rio de Janeiro and Niter[] Brazil: study protocol for a controlled interrupted time series analysis using routine disease surveillance data. <i>F1000Research</i> , 2019 , 8, 1328 Large-Scale Deployment and Establishment of Into the Population in Rio de Janeiro, Brazil. <i>Frontiers in Microbiology</i> , 2021 , 12, 711107 Wolbachia infection does not alter attraction of the mosquito Aedes (Stegomyia) aegypti to human	3.6 5.7	6
29 28 27	Aedes aegypti (Diptera: Culicidae) mosquitoes. <i>Journal of Medical Entomology</i> , 2010 , 47, 1044-52 The impact of large-scale deployment of mosquitoes on arboviral disease incidence in Rio de Janeiro and Niter[] Brazil: study protocol for a controlled interrupted time series analysis using routine disease surveillance data. <i>F1000Research</i> , 2019 , 8, 1328 Large-Scale Deployment and Establishment of Into the Population in Rio de Janeiro, Brazil. <i>Frontiers in Microbiology</i> , 2021 , 12, 711107 Wolbachia infection does not alter attraction of the mosquito Aedes (Stegomyia) aegypti to human odours. <i>Medical and Veterinary Entomology</i> , 2014 , 28, 457-60 In vitro rearing of Perkinsiella saccharicida and the use of leaf segments to assay Fiji disease virus	3.6 5.7 2.4	6 6 5
29 28 27 26	Aedes aegypti (Diptera: Culicidae) mosquitoes. <i>Journal of Medical Entomology</i> , 2010 , 47, 1044-52 The impact of large-scale deployment of mosquitoes on arboviral disease incidence in Rio de Janeiro and Niterij Brazil: study protocol for a controlled interrupted time series analysis using routine disease surveillance data. <i>F1000Research</i> , 2019 , 8, 1328 Large-Scale Deployment and Establishment of Into the Population in Rio de Janeiro, Brazil. <i>Frontiers in Microbiology</i> , 2021 , 12, 711107 Wolbachia infection does not alter attraction of the mosquito Aedes (Stegomyia) aegypti to human odours. <i>Medical and Veterinary Entomology</i> , 2014 , 28, 457-60 In vitro rearing of Perkinsiella saccharicida and the use of leaf segments to assay Fiji disease virus transmission. <i>Phytopathology</i> , 2008 , 98, 810-4	3.6 5.7 2.4	655

22	Semliki Forest virus as an expression vector in insect cell lines. <i>Insect Molecular Biology</i> , 1999 , 8, 409-14	3.4	3
21	Prospects for control of African trypanosomiasis by tsetse vector manipulation. <i>Parasitology Today</i> , 2001 , 17, 29-35		3
20	Response to: Comment on Rohrscheib et al. 2016 "Intensity of mutualism breakdown is determined by temperature not amplification of Wolbachia genes". <i>PLoS Pathogens</i> , 2017 , 13, e1006521	7.6	3
19	Effectiveness of Wolbachia-infected mosquito deployments in reducing the incidence of dengue and other Aedes-borne diseases in Niter[] Brazil: a quasi-experimental study		3
18	Male-Killing Wolbachia in the Butterfly Hypolimnas bolina 2010 , 209-227		2
17	Wolbachia: invasion biology in South pacific butterflies. <i>Current Biology</i> , 2007 , 17, R220-1	6.3	2
16	Crystallization and preliminary diffraction analysis of a DsbA homologue from Wolbachia pipientis. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2008 , 64, 94-7		2
15	Wolbachia: why these bacteria are important to genome research. <i>Microbial & Comparative Genomics</i> , 1999 , 4, 159		2
14	Gene organization of the dnaA region of Wolbachia. <i>Journal of Bacteriology</i> , 1999 , 181, 4708-10	3.5	2
13	Novel phenotype of Wolbachia strain wPip in Aedes aegypti challenges assumptions on mechanisms of Wolbachia-mediated dengue virus inhibition		2
12	Reduced dengue incidence following deployments of Wolbachia-infected Aedes aegypti in Yogyakarta, Indonesia: a quasi-experimental trial using controlled interrupted time series analysis		2
11	The Metabolic Response to Infection With Wolbachia Implicates the Insulin/Insulin-Like-Growth Factor and Hypoxia Signaling Pathways in Drosophila melanogaster. <i>Frontiers in Ecology and Evolution</i> , 2021 , 9,	3.7	2
10	The impact of city-wide deployment of -carrying mosquitoes on arboviral disease incidence in Medelli and Bello, Colombia: study protocol for an interrupted time-series analysis and a test-negative design study <i>F1000Research</i> , 2019 , 8, 1327	3.6	1
9	The impact of city-wide deployment of Wolbachia-carrying mosquitoes on arboviral disease incidence in Medell and Bello, Colombia: study protocol for an interrupted time-series analysis and a test-negative design study. <i>F1000Research</i> ,8, 1327	3.6	1
8	Large-scale deployment and establishment of Wolbachia into the Aedes aegypti population in Rio de Janeiro, Brazil		1
7	Mel genome remains stable after 7 years in Australian field populations. <i>Microbial Genomics</i> , 2021 , 7,	4.4	1
6	Environmental factors influence the local establishment of Wolbachia in Aedes aegypti mosquitoes in two small communities in central Vietnam. <i>Gates Open Research</i> ,5, 147	2.4	1
5	Environmental factors influence the local establishment of Wolbachia in Aedes aegypti mosquitoes	2.4	

LIST OF PUBLICATIONS

4	Aedes aegypti abundance and insecticide resistance profiles in the applying Wolbachia to eliminate dengue trial <i>PLoS Neglected Tropical Diseases</i> , 2022 , 16, e0010284	4.8	1
3	PCR-based detection and identification of insect symbionts 1997 , 561-566		
2	Detection and Identification of Strains in Mosquito Eggs Using Attenuated Total Reflection Fourier Transform Infrared (ATR FT-IR) Spectroscopy. <i>Applied Spectroscopy</i> , 2021 , 75, 1003-1011	3.1	
1	Transient Introgression of Wolbachia into Aedes aegypti Populations Does Not Elicit an Antibody Response to Wolbachia Surface Protein in Community Members. <i>Pathogens</i> , 2022 , 11, 535	4.5	