

George Dimopoulos

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170
papers

14,778
citations

61
h-index

120
g-index

203
ext. papers

17,430
ext. citations

8.1
avg, IF

6.48
L-index

#	Paper	IF	Citations
170	Genome sequence of <i>Aedes aegypti</i> , a major arbovirus vector. <i>Science</i> , 2007 , 316, 1718-23	33.3	867
169	Immunity-related genes and gene families in <i>Anopheles gambiae</i> . <i>Science</i> , 2002 , 298, 159-65	33.3	743
168	The <i>Aedes aegypti</i> toll pathway controls dengue virus infection. <i>PLoS Pathogens</i> , 2008 , 4, e1000098	7.6	578
167	Implication of the mosquito midgut microbiota in the defense against malaria parasites. <i>PLoS Pathogens</i> , 2009 , 5, e1000423	7.6	482
166	Evolutionary dynamics of immune-related genes and pathways in disease-vector mosquitoes. <i>Science</i> , 2007 , 316, 1738-43	33.3	461
165	Comparative genome and proteome analysis of <i>Anopheles gambiae</i> and <i>Drosophila melanogaster</i> . <i>Science</i> , 2002 , 298, 149-59	33.3	455
164	An evolutionary conserved function of the JAK-STAT pathway in anti-dengue defense. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 17841-6	11.5	380
163	Natural microbe-mediated refractoriness to <i>Plasmodium</i> infection in <i>Anopheles gambiae</i> . <i>Science</i> , 2011 , 332, 855-8	33.3	374
162	Sequencing of <i>Culex quinquefasciatus</i> establishes a platform for mosquito comparative genomics. <i>Science</i> , 2010 , 330, 86-8	33.3	352
161	AgDscam, a hypervariable immunoglobulin domain-containing receptor of the <i>Anopheles gambiae</i> innate immune system. <i>PLoS Biology</i> , 2006 , 4, e229	9.7	345
160	Regulation of sexual development of <i>Plasmodium</i> by translational repression. <i>Science</i> , 2006 , 313, 667-9	33.3	333
159	<i>Anopheles gambiae</i> immune responses to human and rodent <i>Plasmodium</i> parasite species. <i>PLoS Pathogens</i> , 2006 , 2, e52	7.6	329
158	<i>Wolbachia</i> invades <i>Anopheles stephensi</i> populations and induces refractoriness to <i>Plasmodium</i> infection. <i>Science</i> , 2013 , 340, 748-51	33.3	307
157	Discovery of insect and human dengue virus host factors. <i>Nature</i> , 2009 , 458, 1047-50	50.4	301
156	Molecular immune responses of the mosquito <i>Anopheles gambiae</i> to bacteria and malaria parasites. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1997 , 94, 11508-13	11.5	301
155	The role of reactive oxygen species on <i>Plasmodium melanotic</i> encapsulation in <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 14139-44	11.5	237
154	Genome expression analysis of <i>Anopheles gambiae</i> : responses to injury, bacterial challenge, and malaria infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002 , 99, 8814-9	11.5	236

153	Malaria infection of the mosquito <i>Anopheles gambiae</i> activates immune-responsive genes during critical transition stages of the parasite life cycle. <i>EMBO Journal</i> , 1998 , 17, 6115-23	13	230
152	Genome of <i>Rhodnius prolixus</i> , an insect vector of Chagas disease, reveals unique adaptations to hematophagy and parasite infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 14936-41	11.5	220
151	CTRP is essential for mosquito infection by malaria ookinetes. <i>EMBO Journal</i> , 1999 , 18, 6221-7	13	219
150	Reciprocal tripartite interactions between the <i>Aedes aegypti</i> midgut microbiota, innate immune system and dengue virus influences vector competence. <i>PLoS Neglected Tropical Diseases</i> , 2012 , 6, e1564.8	4.8	215
149	A hemocyte-like cell line established from the malaria vector <i>Anopheles gambiae</i> expresses six prophenoloxidase genes. <i>Journal of Biological Chemistry</i> , 1999 , 274, 11727-35	5.4	206
148	Universal features of post-transcriptional gene regulation are critical for <i>Plasmodium</i> zygote development. <i>PLoS Pathogens</i> , 2010 , 6, e1000767	7.6	180
147	Native microbiota shape insect vector competence for human pathogens. <i>Cell Host and Microbe</i> , 2011 , 10, 307-10	23.4	168
146	Caspar controls resistance to <i>Plasmodium falciparum</i> in diverse anopheline species. <i>PLoS Pathogens</i> , 2009 , 5, e1000335	7.6	165
145	Gambicin: a novel immune responsive antimicrobial peptide from the malaria vector <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001 , 98, 12630-5	11.5	153
144	P25 and P28 proteins of the malaria ookinete surface have multiple and partially redundant functions. <i>EMBO Journal</i> , 2001 , 20, 3975-83	13	152
143	Chromobacterium Csp_P reduces malaria and dengue infection in vector mosquitoes and has entomopathogenic and in vitro anti-pathogen activities. <i>PLoS Pathogens</i> , 2014 , 10, e1004398	7.6	151
142	<i>Anopheles</i> fibrinogen-related proteins provide expanded pattern recognition capacity against bacteria and malaria parasites. <i>Journal of Biological Chemistry</i> , 2009 , 284, 9835-44	5.4	147
141	<i>Plasmodium</i> activates the innate immune response of <i>Anopheles gambiae</i> mosquitoes. <i>EMBO Journal</i> , 1997 , 16, 6114-9	13	147
140	Mosquito immune defenses against <i>Plasmodium</i> infection. <i>Developmental and Comparative Immunology</i> , 2010 , 34, 387-95	3.2	145
139	Dengue virus infection of the <i>Aedes aegypti</i> salivary gland and chemosensory apparatus induces genes that modulate infection and blood-feeding behavior. <i>PLoS Pathogens</i> , 2012 , 8, e1002631	7.6	138
138	The Toll immune signaling pathway control conserved anti-dengue defenses across diverse <i>Ae. aegypti</i> strains and against multiple dengue virus serotypes. <i>Developmental and Comparative Immunology</i> , 2010 , 34, 625-9	3.2	134
137	<i>Anopheles gambiae</i> pilot gene discovery project: identification of mosquito innate immunity genes from expressed sequence tags generated from immune-competent cell lines. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000 , 97, 6619-24	11.5	132
136	Trapping cDNAs encoding secreted proteins from the salivary glands of the malaria vector <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999 , 96, 1516-21	11.5	132

135	Engineered anopheles immunity to Plasmodium infection. <i>PLoS Pathogens</i> , 2011 , 7, e1002458	7.6	130
134	The mosquito microbiota influences vector competence for human pathogens. <i>Current Opinion in Insect Science</i> , 2014 , 3, 6-13	5.1	126
133	Insect immunity and its implication in mosquito-malaria interactions. <i>Cellular Microbiology</i> , 2003 , 5, 3-14	3.9	125
132	Cloning and analysis of a cecropin gene from the malaria vector mosquito, <i>Anopheles gambiae</i> . <i>Insect Molecular Biology</i> , 2000 , 9, 75-84	3.4	124
131	Pathogenomics of <i>Culex quinquefasciatus</i> and meta-analysis of infection responses to diverse pathogens. <i>Science</i> , 2010 , 330, 88-90	33.3	120
130	Dengue virus inhibits immune responses in <i>Aedes aegypti</i> cells. <i>PLoS ONE</i> , 2010 , 5, e10678	3.7	115
129	SOAP, a novel malaria ookinete protein involved in mosquito midgut invasion and oocyst development. <i>Molecular Microbiology</i> , 2003 , 49, 319-29	4.1	107
128	Innate immune defense against malaria infection in the mosquito. <i>Current Opinion in Immunology</i> , 2001 , 13, 79-88	7.8	107
127	The insect microbiome modulates vector competence for arboviruses. <i>Viruses</i> , 2014 , 6, 4294-313	6.2	105
126	Transcriptomic profiling of diverse <i>Aedes aegypti</i> strains reveals increased basal-level immune activation in dengue virus-refractory populations and identifies novel virus-vector molecular interactions. <i>PLoS Neglected Tropical Diseases</i> , 2013 , 7, e2295	4.8	105
125	Mosquito immunity against arboviruses. <i>Viruses</i> , 2014 , 6, 4479-504	6.2	102
124	Exploring <i>Anopheles</i> gut bacteria for <i>Plasmodium</i> blocking activity. <i>Environmental Microbiology</i> , 2014 , 16, 2980-94	5.2	101
123	The <i>Anopheles</i> innate immune system in the defense against malaria infection. <i>Journal of Innate Immunity</i> , 2014 , 6, 169-81	6.9	98
122	<i>Anopheles</i> Imd pathway factors and effectors in infection intensity-dependent anti- <i>Plasmodium</i> action. <i>PLoS Pathogens</i> , 2012 , 8, e1002737	7.6	92
121	Global gene expression analysis of <i>Anopheles gambiae</i> responses to microbial challenge. <i>Insect Biochemistry and Molecular Biology</i> , 2005 , 35, 709-19	4.5	86
120	Genome-wide transcriptomic profiling of <i>Anopheles gambiae</i> hemocytes reveals pathogen-specific signatures upon bacterial challenge and <i>Plasmodium berghei</i> infection. <i>BMC Genomics</i> , 2009 , 10, 257	4.5	83
119	Engineered <i>Aedes aegypti</i> JAK/STAT Pathway-Mediated Immunity to Dengue Virus. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005187	4.8	81
118	CRISPR/Cas9 -mediated gene knockout of <i>Anopheles gambiae</i> FREP1 suppresses malaria parasite infection. <i>PLoS Pathogens</i> , 2018 , 14, e1006898	7.6	80

117	Identification and characterization of differentially expressed cDNAs of the vector mosquito, <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996 , 93, 13066-71	11.5	79
116	<i>Anopheles</i> NF- κ B-regulated splicing factors direct pathogen-specific repertoires of the hypervariable pattern recognition receptor AgDscam. <i>Cell Host and Microbe</i> , 2012 , 12, 521-30	23.4	77
115	The entomopathogenic fungus <i>Beauveria bassiana</i> activate toll and JAK-STAT pathway-controlled effector genes and anti-dengue activity in <i>Aedes aegypti</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2012 , 42, 126-32	4.5	74
114	Transcriptome analysis of <i>Aedes aegypti</i> transgenic mosquitoes with altered immunity. <i>PLoS Pathogens</i> , 2011 , 7, e1002394	7.6	69
113	Analysis of the <i>Plasmodium</i> and <i>Anopheles</i> transcriptional repertoire during ookinete development and midgut invasion. <i>Journal of Biological Chemistry</i> , 2004 , 279, 5573-80	5.4	69
112	Molecular Responses to Zika Virus: Modulation of Infection by the Toll and Jak/Stat Immune Pathways and Virus Host Factors. <i>Frontiers in Microbiology</i> , 2017 , 8, 2050	5.7	64
111	The Gram-negative bacteria-binding protein gene family: its role in the innate immune system of <i>Anopheles gambiae</i> and in anti- <i>Plasmodium</i> defence. <i>Insect Molecular Biology</i> , 2008 , 17, 39-51	3.4	62
110	Transcript profiles of <i>Blumeria graminis</i> development during infection reveal a cluster of genes that are potential virulence determinants. <i>Molecular Plant-Microbe Interactions</i> , 2005 , 18, 125-33	3.6	62
109	Analysis of the <i>Plasmodium</i> and <i>Anopheles</i> transcriptomes during oocyst differentiation. <i>Journal of Biological Chemistry</i> , 2004 , 279, 5581-7	5.4	61
108	Mosquito gut antiparasitic and antiviral immunity. <i>Developmental and Comparative Immunology</i> , 2016 , 64, 53-64	3.2	57
107	A cell surface mucin specifically expressed in the midgut of the malaria mosquito <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999 , 96, 5610-5	11.5	57
106	Transcriptomic and functional analysis of the <i>Anopheles gambiae</i> salivary gland in relation to blood feeding. <i>BMC Genomics</i> , 2010 , 11, 566	4.5	56
105	Mosquito infection responses to developing filarial worms. <i>PLoS Neglected Tropical Diseases</i> , 2009 , 3, e529	4.8	53
104	Gene discovery in an invasive tephritid model pest species, the Mediterranean fruit fly, <i>Ceratitidis capitata</i> . <i>BMC Genomics</i> , 2008 , 9, 243	4.5	53
103	Transcriptome profiling of sexual maturation and mating in the Mediterranean fruit fly, <i>Ceratitidis capitata</i> . <i>PLoS ONE</i> , 2012 , 7, e30857	3.7	53
102	Transcriptome analysis of <i>Anopheles stephensi</i> - <i>Plasmodium berghei</i> interactions. <i>Molecular and Biochemical Parasitology</i> , 2005 , 142, 76-87	1.9	51
101	Emerging role of lipid droplets in <i>Aedes aegypti</i> immune response against bacteria and Dengue virus. <i>Scientific Reports</i> , 2016 , 6, 19928	4.9	49
100	Polymorphisms detected by random PCR distinguish between different chromosomal forms of <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1994 , 91, 10315-9	11.5	49

99	MicroRNA-regulation of <i>Anopheles gambiae</i> immunity to <i>Plasmodium falciparum</i> infection and midgut microbiota. <i>Developmental and Comparative Immunology</i> , 2015 , 49, 170-8	3.2	48
98	Immunoglobulin superfamily members play an important role in the mosquito immune system. <i>Developmental and Comparative Immunology</i> , 2008 , 32, 519-31	3.2	47
97	Bayesian coclustering of <i>Anopheles</i> gene expression time series: study of immune defense response to multiple experimental challenges. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 16939-44	11.5	46
96	<i>Aedes aegypti</i> ML and Niemann-Pick type C family members are agonists of dengue virus infection. <i>Developmental and Comparative Immunology</i> , 2014 , 43, 1-9	3.2	45
95	An -associated fungus increases susceptibility to dengue virus by modulating gut trypsin activity. <i>ELife</i> , 2017 , 6,	8.9	45
94	Molecular analysis of photic inhibition of blood-feeding in <i>Anopheles gambiae</i> . <i>BMC Physiology</i> , 2008 , 8, 23	0	43
93	Changes in the microbiota cause genetically modified to spread in a population. <i>Science</i> , 2017 , 357, 1396-1399	3.3	40
92	Integrated genetic map of <i>Anopheles gambiae</i> : use of RAPD polymorphisms for genetic, cytogenetic and STS landmarks. <i>Genetics</i> , 1996 , 143, 953-60	4	40
91	<i>Anopheles</i> Midgut FREP1 Mediates <i>Plasmodium</i> Invasion. <i>Journal of Biological Chemistry</i> , 2015 , 290, 16490-50139	3.4	39
90	Cytoplasmic actin is an extracellular insect immune factor which is secreted upon immune challenge and mediates phagocytosis and direct killing of bacteria, and is a <i>Plasmodium</i> Antagonist. <i>PLoS Pathogens</i> , 2015 , 11, e1004631	7.6	39
89	Gametocytocidal screen identifies novel chemical classes with <i>Plasmodium falciparum</i> transmission blocking activity. <i>PLoS ONE</i> , 2014 , 9, e105817	3.7	39
88	<i>Anopheles</i> infection responses; laboratory models versus field malaria transmission systems. <i>Acta Tropica</i> , 2005 , 95, 285-91	3.2	38
87	Challenges and approaches for mosquito targeted malaria control. <i>Current Molecular Medicine</i> , 2009 , 9, 116-30	2.5	37
86	Heme Signaling Impacts Global Gene Expression, Immunity and Dengue Virus Infectivity in <i>Aedes aegypti</i> . <i>PLoS ONE</i> , 2015 , 10, e0135985	3.7	35
85	Suppressing dengue-2 infection by chemical inhibition of <i>Aedes aegypti</i> host factors. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e3084	4.8	34
84	Regulation of midgut cell proliferation impacts <i>Aedes aegypti</i> susceptibility to dengue virus. <i>PLoS Neglected Tropical Diseases</i> , 2018 , 12, e0006498	4.8	34
83	Caudal is a negative regulator of the <i>Anopheles</i> IMD pathway that controls resistance to <i>Plasmodium falciparum</i> infection. <i>Developmental and Comparative Immunology</i> , 2013 , 39, 323-32	3.2	33
82	Spatial and sex-specific dissection of the <i>Anopheles gambiae</i> midgut transcriptome. <i>BMC Genomics</i> , 2007 , 8, 37	4.5	33

81	The effect of hyperosmotic pressure on antibody production and gene expression in the GS-NSO cell line. <i>Biotechnology and Applied Biochemistry</i> , 2004 , 40, 41-6	2.8	31
80	Analysis of the <i>Anopheles gambiae</i> genome using RAPD markers. <i>Insect Molecular Biology</i> , 1994 , 3, 149-574	3.4	31
79	Protocol for mosquito rearing (<i>A. gambiae</i>). <i>Journal of Visualized Experiments</i> , 2007 , 221	1.6	30
78	<i>Anopheles gambiae</i> immune responses to Sephadex beads: involvement of anti-Plasmodium factors in regulating melanization. <i>Insect Biochemistry and Molecular Biology</i> , 2006 , 36, 769-78	4.5	29
77	Amino acid metabolic signaling influences <i>Aedes aegypti</i> midgut microbiome variability. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005677	4.8	29
76	Functional genomic analyses of <i>Enterobacter</i> , <i>Anopheles</i> and <i>Plasmodium</i> reciprocal interactions that impact vector competence. <i>Malaria Journal</i> , 2016 , 15, 425	3.6	29
75	Bacteria- and IMD pathway-independent immune defenses against <i>Plasmodium falciparum</i> in <i>Anopheles gambiae</i> . <i>PLoS ONE</i> , 2013 , 8, e72130	3.7	28
74	Cloning and molecular characterization of two mosquito iron regulatory proteins. <i>Insect Biochemistry and Molecular Biology</i> , 2002 , 32, 579-89	4.5	27
73	The Golgi associated ERI3 is a Flavivirus host factor. <i>Scientific Reports</i> , 2016 , 6, 34379	4.9	27
72	Curious entanglements: interactions between mosquitoes, their microbiota, and arboviruses. <i>Current Opinion in Virology</i> , 2019 , 37, 26-36	7.5	26
71	<i>Chromobacterium</i> spp. mediate their anti-Plasmodium activity through secretion of the histone deacetylase inhibitor romidepsin. <i>Scientific Reports</i> , 2018 , 8, 6176	4.9	25
70	Diverse Host and Restriction Factors Regulate Mosquito-Pathogen Interactions. <i>Trends in Parasitology</i> , 2018 , 34, 603-616	6.4	25
69	Aminopeptidase secreted by <i>Chromobacterium</i> sp. Panama inhibits dengue virus infection by degrading the E protein. <i>PLoS Neglected Tropical Diseases</i> , 2018 , 12, e0006443	4.8	24
68	A natural <i>Anopheles</i> -associated <i>Penicillium chrysogenum</i> enhances mosquito susceptibility to <i>Plasmodium</i> infection. <i>Scientific Reports</i> , 2016 , 6, 34084	4.9	23
67	The <i>Anopheles</i> FBN9 immune factor mediates <i>Plasmodium</i> species-specific defense through transgenic fat body expression. <i>Developmental and Comparative Immunology</i> , 2017 , 67, 257-265	3.2	23
66	Protocol for dengue infections in mosquitoes (<i>A. aegypti</i>) and infection phenotype determination. <i>Journal of Visualized Experiments</i> , 2007 , 220	1.6	23
65	Dual roles for the ER membrane protein complex in flavivirus infection: viral entry and protein biogenesis. <i>Scientific Reports</i> , 2019 , 9, 9711	4.9	22
64	Immune Regulation of Is Species Specific and Infection Intensity Dependent. <i>MBio</i> , 2017 , 8,	7.8	22

63	Influences of the Mosquito Microbiota on Vector Competence. <i>Advances in Insect Physiology</i> , 2016 , 51, 243-291	2.5	22
62	Versatile transgenic multistage effector-gene combinations for suppression in. <i>Science Advances</i> , 2020 , 6, eaay5898	14.3	21
61	A mosquito salivary gland protein partially inhibits Plasmodium sporozoite cell traversal and transmission. <i>Nature Communications</i> , 2018 , 9, 2908	17.4	21
60	Salivary gland-specific gene expression in the malaria vector <i>Anopheles gambiae</i> . <i>Parassitologia</i> , 1999 , 41, 483-7		20
59	Immunization with AgTRIO, a Protein in <i>Anopheles Saliva</i> , Contributes to Protection against Plasmodium Infection in Mice. <i>Cell Host and Microbe</i> , 2018 , 23, 523-535.e5	23.4	18
58	Identification of anti-flaviviral drugs with mosquitocidal and anti-Zika virus activity in <i>Aedes aegypti</i> . <i>PLoS Neglected Tropical Diseases</i> , 2019 , 13, e0007681	4.8	18
57	Protocol for RNAi assays in adult mosquitoes (<i>A. gambiae</i>). <i>Journal of Visualized Experiments</i> , 2007 , 230	1.6	18
56	Low- and high-tech approaches to control Plasmodium parasite transmission by anopheles mosquitoes. <i>Journal of Tropical Medicine</i> , 2011 , 2011, 891342	2.4	17
55	Genome-wide analysis of transcriptomic divergence between laboratory colony and field <i>Anopheles gambiae</i> mosquitoes of the M and S molecular forms. <i>Insect Molecular Biology</i> , 2010 , 19, 695-705	3.4	17
54	Comprehensive DNA methylation analysis of the <i>Aedes aegypti</i> genome. <i>Scientific Reports</i> , 2016 , 6, 36444	4.9	16
53	Association of microRNAs with Argonaute proteins in the malaria mosquito <i>Anopheles gambiae</i> after blood ingestion. <i>Scientific Reports</i> , 2017 , 7, 6493	4.9	16
52	Chloroquine mediated modulation of <i>Anopheles gambiae</i> gene expression. <i>PLoS ONE</i> , 2008 , 3, e2587	3.7	15
51	Characterization of the Rel2-regulated transcriptome and proteome of <i>Anopheles stephensi</i> identifies new anti-Plasmodium factors. <i>Insect Biochemistry and Molecular Biology</i> , 2014 , 52, 82-93	4.5	14
50	Functional genomics studies on the innate immunity of disease vectors. <i>Insect Science</i> , 2008 , 15, 15-27	3.6	14
49	Broad spectrum immunomodulatory effects of <i>Anopheles gambiae</i> microRNAs and their use for transgenic suppression of Plasmodium. <i>PLoS Pathogens</i> , 2020 , 16, e1008453	7.6	13
48	Disruption of mosGILT in <i>Anopheles gambiae</i> impairs ovarian development and Plasmodium infection. <i>Journal of Experimental Medicine</i> , 2020 , 217,	16.6	12
47	Molecular discrimination of mosquito vectors and their pathogens. <i>Expert Review of Molecular Diagnostics</i> , 2009 , 9, 757-65	3.8	11
46	New rapid one-step PCR diagnostic assay for Plasmodium falciparum infective mosquitoes. <i>Scientific Reports</i> , 2018 , 8, 1462	4.9	10

45	Malaria parasites and the anopheles mosquito. <i>Chemical Immunology and Allergy</i> , 2002 , 80, 27-49		10
44	A mosquito mediator of parasite-induced immune priming. <i>Trends in Parasitology</i> , 2015 , 31, 402-4	6.4	9
43	Larval exposure to bacteria modulates arbovirus infection and immune gene expression in adult <i>Aedes aegypti</i> . <i>Developmental and Comparative Immunology</i> , 2020 , 104, 103540	3.2	9
42	Prospects and Pitfalls: Next-Generation Tools to Control Mosquito-Transmitted Disease. <i>Annual Review of Microbiology</i> , 2020 , 74, 455-475	17.5	9
41	Hydrogen cyanide produced by the soil bacterium <i>Chromobacterium</i> sp. Panama contributes to mortality in <i>Anopheles gambiae</i> mosquito larvae. <i>Scientific Reports</i> , 2018 , 8, 8358	4.9	9
40	Dynamic miRNA-mRNA interactions coordinate gene expression in adult <i>Anopheles gambiae</i> . <i>PLoS Genetics</i> , 2020 , 16, e1008765	6	8
39	Continuous exposure to <i>Plasmodium</i> results in decreased susceptibility and transcriptomic divergence of the <i>Anopheles gambiae</i> immune system. <i>BMC Genomics</i> , 2007 , 8, 451	4.5	8
38	Inhibition of in Adult Mosquitoes Causes Male-Specific Mortality and Diverse Transcriptome Changes. <i>Pathogens</i> , 2020 , 9,	4.5	7
37	Molecular analysis of <i>Culex quinquefasciatus</i> larvae responses to <i>Lysinibacillus sphaericus</i> Bin toxin. <i>PLoS ONE</i> , 2017 , 12, e0175473	3.7	7
36	<i>Plasmodium falciparum</i> Gametocyte Culture and Mosquito Infection Through Artificial Membrane Feeding. <i>Journal of Visualized Experiments</i> , 2020 ,	1.6	7
35	Application of the Relationship-Based Model to Engagement for Field Trials of Genetically Engineered Malaria Vectors. <i>American Journal of Tropical Medicine and Hygiene</i> , 2020 ,	3.2	7
34	Bacterial natural products in the fight against mosquito-transmitted tropical diseases. <i>Natural Product Reports</i> , 2020 , 37, 338-354	15.1	7
33	Glyphosate inhibits melanization and increases susceptibility to infection in insects. <i>PLoS Biology</i> , 2021 , 19, e3001182	9.7	7
32	Mosquito antiviral immune pathways. <i>Developmental and Comparative Immunology</i> , 2021 , 116, 103964	3.2	7
31	The mosquito adulticidal <i>Chromobacterium</i> sp. Panama causes transgenerational impacts on fitness parameters and elicits xenobiotic gene responses. <i>Parasites and Vectors</i> , 2018 , 11, 229	4	6
30	Transcriptional mediators Kto and Skd are involved in the regulation of the IMD pathway and anti- <i>Plasmodium</i> defense in <i>Anopheles gambiae</i> . <i>PLoS ONE</i> , 2012 , 7, e45580	3.7	6
29	Immune response-related genes associated to blocking midgut dengue virus infection in <i>Aedes aegypti</i> strains that differ in susceptibility. <i>Insect Science</i> , 2019 , 26, 635-648	3.6	6
28	How does <i>Anopheles gambiae</i> kill malaria parasites?. <i>Parassitologia</i> , 1999 , 41, 169-75		6

27	Glucose-mediated proliferation of a gut commensal bacterium promotes Plasmodium infection by increasing mosquito midgut pH. <i>Cell Reports</i> , 2021 , 35, 108992	10.6	5
26	Microbial Diversity of Adult Aedes aegypti and Water Collected from Different Mosquito Aquatic Habitats in Puerto Rico. <i>Microbial Ecology</i> , 2021 , 1	4.4	5
25	Acoustotactic response of mosquitoes in untethered flight to incidental sound. <i>Scientific Reports</i> , 2021 , 11, 1884	4.9	5
24	A Nonlive Preparation of sp. Panama (Csp_P) Is a Highly Effective Larval Mosquito Biopesticide. <i>Applied and Environmental Microbiology</i> , 2020 , 86,	4.8	4
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8	Field-deployable molecular diagnostic platform for arbovirus detection in <i>Aedes aegypti</i> . <i>Parasites and Vectors</i> , 2020 , 13, 489	4	1
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