

Elena A Levashina

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9386467/publications.pdf>

Version: 2024-02-01

76
papers

7,323
citations

71061

41
h-index

91828

69
g-index

87
all docs

87
docs citations

87
times ranked

6316
citing authors

#	ARTICLE	IF	CITATIONS
1	A high-affinity antibody against the CSP N-terminal domain lacks inhibitory activity. <i>Journal of Experimental Medicine</i> , 2020, 217, .	4.2	21
2	Kinetics of Plasmodium midgut invasion in Anopheles mosquitoes. <i>PLoS Pathogens</i> , 2020, 16, e1008739.	2.1	12
3	Evolution of protective human antibodies against Plasmodium falciparum circumsporozoite protein repeat motifs. <i>Nature Medicine</i> , 2020, 26, 1135-1145.	15.2	64
4	Critical Steps of Plasmodium falciparum Ookinete Maturation. <i>Frontiers in Microbiology</i> , 2020, 11, 269.	1.5	22
5	Kinetics of Plasmodium midgut invasion in Anopheles mosquitoes. , 2020, 16, e1008739.		0
6	Kinetics of Plasmodium midgut invasion in Anopheles mosquitoes. , 2020, 16, e1008739.		0
7	Kinetics of Plasmodium midgut invasion in Anopheles mosquitoes. , 2020, 16, e1008739.		0
8	Kinetics of Plasmodium midgut invasion in Anopheles mosquitoes. , 2020, 16, e1008739.		0
9	Anopheles gambiae TEP1 forms a complex with the coiled-coil domain of LRIM1/APL1C following a conformational change in the thioester domain. <i>PLoS ONE</i> , 2019, 14, e0218203.	1.1	24
10	Mosquito microevolution drives Plasmodium falciparum dynamics. <i>Nature Microbiology</i> , 2019, 4, 941-947.	5.9	18
11	A Novel Tool for the Generation of Conditional Knockouts To Study Gene Function across the Plasmodium falciparum Life Cycle. <i>MBio</i> , 2019, 10, .	1.8	45
12	Metabolic balancing by miR-276 shapes the mosquito reproductive cycle and Plasmodium falciparum development. <i>Nature Communications</i> , 2019, 10, 5634.	5.8	31
13	Clonal selection drives protective memory B cell responses in controlled human malaria infection. <i>Science Immunology</i> , 2018, 3, .	5.6	132
14	MicroRNA Tissue Atlas of the Malaria Mosquito <i>Anopheles gambiae</i> . <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 185-193.	0.8	21
15	Rare PfCSP C-terminal antibodies induced by live sporozoite vaccination are ineffective against malaria infection. <i>Journal of Experimental Medicine</i> , 2018, 215, 63-75.	4.2	79
16	Vector Immunity and Evolutionary Ecology: The Harmonious Dissonance. <i>Trends in Immunology</i> , 2018, 39, 862-873.	2.9	33
17	Micromanaging Immunity in the Murine Host vs. the Mosquito Vector: Microbiota-Dependent Immune Responses to Intestinal Parasites. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018, 8, 308.	1.8	10
18	Non-competitive resource exploitation within mosquito shapes within-host malaria infectivity and virulence. <i>Nature Communications</i> , 2018, 9, 3474.	5.8	58

#	ARTICLE	IF	CITATIONS
19	Complement-Like System in the Mosquito Responses Against Malaria Parasites. , 2018, , 139-146.		2
20	Unbiased classification of mosquito blood cells by single-cell genomics and high-content imaging. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E7568-E7577.	3.3	57
21	Insects Go on a STING Operation to Tackle Intracellular Invaders. Immunity, 2018, 49, 195-197.	6.6	4
22	Antihomotypic affinity maturation improves human B cell responses against a repetitive epitope. Science, 2018, 360, 1358-1362.	6.0	89
23	The role of microRNA in <i>Anopheles</i> biology—an emerging research field. Parasite Immunology, 2017, 39, e12405.	0.7	11
24	Natural Parasite Exposure Induces Protective Human Anti-Malarial Antibodies. Immunity, 2017, 47, 1197-1209.e10.	6.6	129
25	NF- κ B-Like Signaling Pathway REL2 in Immune Defenses of the Malaria Vector <i>Anopheles gambiae</i> . Frontiers in Cellular and Infection Microbiology, 2017, 7, 258.	1.8	18
26	Variation in susceptibility of African <i>Plasmodium falciparum</i> malaria parasites to TEP1 mediated killing in <i>Anopheles gambiae</i> mosquitoes. Scientific Reports, 2016, 6, 20440.	1.6	34
27	Comparative Proteomics and Functional Analysis Reveal a Role of <i>Plasmodium falciparum</i> Osmiophilic Bodies in Malaria Parasite Transmission. Molecular and Cellular Proteomics, 2016, 15, 3243-3255.	2.5	40
28	Advancing vector biology research: a community survey for future directions, research applications and infrastructure requirements. Pathogens and Global Health, 2016, 110, 164-172.	1.0	3
29	The reproductive tracts of two malaria vectors are populated by a core microbiome and by gender- and swarm-enriched microbial biomarkers. Scientific Reports, 2016, 6, 24207.	1.6	93
30	Tools for <i>Anopheles gambiae</i> Transgenesis. G3: Genes, Genomes, Genetics, 2015, 5, 1151-1163.	0.8	95
31	A New Role of the Mosquito Complement-like Cascade in Male Fertility in <i>Anopheles gambiae</i> . PLoS Biology, 2015, 13, e1002255.	2.6	53
32	Mosquito defenses against <i>Plasmodium</i> parasites. Current Opinion in Insect Science, 2014, 3, 30-36.	2.2	18
33	Evidence of natural <i>Wolbachia</i> infections in field populations of <i>Anopheles gambiae</i> . Nature Communications, 2014, 5, 3985.	5.8	142
34	Transcriptome-wide analysis of microRNA expression in the malaria mosquito <i>Anopheles gambiae</i> . BMC Genomics, 2014, 15, 557.	1.2	49
35	AP-1/Fos-TGase2 Axis Mediates Wounding-induced <i>Plasmodium falciparum</i> Killing in <i>Anopheles gambiae</i> . Journal of Biological Chemistry, 2013, 288, 16145-16154.	1.6	22
36	Targeted Mutagenesis in the Malaria Mosquito Using TALE Nucleases. PLoS ONE, 2013, 8, e74511.	1.1	78

#	ARTICLE	IF	CITATIONS
37	Midgut Microbiota of the Malaria Mosquito Vector <i>Anopheles gambiae</i> and Interactions with <i>Plasmodium falciparum</i> Infection. <i>PLoS Pathogens</i> , 2012, 8, e1002742.	2.1	427
38	Silencing of Genes and Alleles by RNAi in <i>Anopheles gambiae</i> . <i>Methods in Molecular Biology</i> , 2012, 923, 161-176.	0.4	8
39	High-throughput sorting of mosquito larvae for laboratory studies and for future vector control interventions. <i>Malaria Journal</i> , 2012, 11, 302.	0.8	56
40	Salivary Gland-Specific <i>P. berghei</i> Reporter Lines Enable Rapid Evaluation of Tissue-Specific Sporozoite Loads in Mosquitoes. <i>PLoS ONE</i> , 2012, 7, e36376.	1.1	15
41	Lawrence's book review unfair to Hoffmann. <i>Current Biology</i> , 2012, 22, R482.	1.8	0
42	Genetic clonality of <i>Plasmodium falciparum</i> affects the outcome of infection in <i>Anopheles gambiae</i> . <i>International Journal for Parasitology</i> , 2012, 42, 589-595.	1.3	44
43	Intracellular immune responses of dipteran insects. <i>Immunological Reviews</i> , 2011, 240, 129-140.	2.8	31
44	Functional Genomics of Tick Thioester-Containing Proteins Reveal the Ancient Origin of the Complement System. <i>Journal of Innate Immunity</i> , 2011, 3, 623-630.	1.8	55
45	Focusing on complement in the antiparasitic defense of mosquitoes. <i>Trends in Parasitology</i> , 2010, 26, 1-3.	1.5	13
46	A heterodimeric complex of the LRR proteins LRIM1 and APL1C regulates complement-like immunity in <i>Anopheles gambiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 16817-16822.	3.3	58
47	The Major Yolk Protein Vitellogenin Interferes with the Anti- <i>Plasmodium</i> Response in the Malaria Mosquito <i>Anopheles gambiae</i> . <i>PLoS Biology</i> , 2010, 8, e1000434.	2.6	144
48	Two Mosquito LRR Proteins Function as Complement Control Factors in the TEP1-Mediated Killing of <i>Plasmodium</i> . <i>Cell Host and Microbe</i> , 2009, 5, 273-284.	5.1	212
49	Dissecting the Genetic Basis of Resistance to Malaria Parasites in <i>Anopheles gambiae</i> . <i>Science</i> , 2009, 326, 147-150.	6.0	106
50	RNAi in the Malaria Vector, <i>Anopheles gambiae</i> . <i>Methods in Molecular Biology</i> , 2009, 555, 63-75.	0.4	17
51	Host-parasite interactions: the balance of trade. <i>Current Opinion in Microbiology</i> , 2008, 11, 338-339.	2.3	0
52	Antimalarial Responses in <i>Anopheles gambiae</i> : From a Complement-like Protein to a Complement-like Pathway. <i>Cell Host and Microbe</i> , 2008, 3, 364-374.	5.1	125
53	Molecular and cellular components of the mating machinery in <i>Anopheles gambiae</i> females. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 19390-19395.	3.3	107
54	Reverse Genetics Analysis of Antiparasitic Responses in the Malaria Vector, <i>Anopheles gambiae</i> . , 2008, 415, 365-377.		6

#	ARTICLE	IF	CITATIONS
55	Structural basis for conserved complement factor-like function in the antimalarial protein TEP1. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 11615-11620.	3.3	87
56	Evolutionary Dynamics of Immune-Related Genes and Pathways in Disease-Vector Mosquitoes. Science, 2007, 316, 1738-1743.	6.0	550
57	Malaria Plasmodium agent induces alteration in the head proteome of their Anopheles mosquito host. Proteomics, 2007, 7, 1908-1915.	1.3	75
58	Phagocytosis in mosquito immune responses. Immunological Reviews, 2007, 219, 8-16.	2.8	53
59	Boosting NF- κ B-Dependent Basal Immunity of <i>Anopheles gambiae</i> Aborts Development of <i>Plasmodium berghei</i> . Immunity, 2006, 25, 677-685.	6.6	210
60	Fz2 and Cdc42 Mediate Melanization and Actin Polymerization but Are Dispensable for Plasmodium Killing in the Mosquito Midgut. PLoS Pathogens, 2006, 2, e133.	2.1	46
61	In Vivo Identification of Novel Regulators and Conserved Pathways of Phagocytosis in <i>A. gambiae</i> . Immunity, 2005, 23, 65-73.	6.6	126
62	Immune responses and parasite transmission in blood-feeding insects. Trends in Parasitology, 2004, 20, 433-439.	1.5	51
63	Mosquito immune responses against malaria parasites. Current Opinion in Immunology, 2004, 16, 16-20.	2.4	48
64	Bacterial alpha2-macroglobulins: colonization factors acquired by horizontal gene transfer from the metazoan genome?. Genome Biology, 2004, 5, R38.	13.9	74
65	Thioester-containing proteins and insect immunity. Molecular Immunology, 2004, 40, 903-908.	1.0	188
66	Immune responses in <i>Anopheles gambiae</i> . Insect Biochemistry and Molecular Biology, 2004, 34, 673-678.	1.2	55
67	Complement-Like Protein TEP1 Is a Determinant of Vectorial Capacity in the Malaria Vector <i>Anopheles gambiae</i> . Cell, 2004, 116, 661-670.	13.5	566
68	Silencing of Toll pathway components by direct injection of double-stranded RNA into <i>Drosophila</i> adult flies. Nucleic Acids Research, 2003, 31, 6619-6623.	6.5	63
69	Immunity-Related Genes and Gene Families in <i>Anopheles gambiae</i> . Science, 2002, 298, 159-165.	6.0	845
70	Reverse genetics in the mosquito <i>Anopheles gambiae</i> : targeted disruption of the Defensin gene. EMBO Reports, 2002, 3, 852-856.	2.0	331
71	Conserved Role of a Complement-like Protein in Phagocytosis Revealed by dsRNA Knockout in Cultured Cells of the Mosquito, <i>Anopheles gambiae</i> . Cell, 2001, 104, 709-718.	13.5	472
72	Innate immune defense against malaria infection in the mosquito. Current Opinion in Immunology, 2001, 13, 79-88.	2.4	116

#	ARTICLE	IF	CITATIONS
73	The <i>necrotic</i> Gene in <i>Drosophila</i> Corresponds to One of a Cluster of Three Serpin Transcripts Mapping at 43A1.2. <i>Genetics</i> , 2000, 156, 1117-1127.	1.2	66
74	Two distinct pathways can control expression of the gene encoding the <i>Drosophila</i> antimicrobial peptide metchnikowin. <i>Journal of Molecular Biology</i> , 1998, 278, 515-527.	2.0	120
75	Similarities between insect and plant host defences. <i>Trends in Cell Biology</i> , 1997, 7, 316.	3.6	0
76	Metchnikowin, a Novel Immune-Inducible Proline-Rich Peptide from <i>Drosophila</i> with Antibacterial and Antifungal Properties. <i>FEBS Journal</i> , 1995, 233, 694-700.	0.2	191