Oliver Billker

List of Publications by Year in descending order

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47006 64796 8,250 86 47 79 citations h-index g-index papers 94 94 94 6581 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Identification of xanthurenic acid as the putative inducer of malaria development in the mosquito. Nature, 1998, 392, 289-292.	27.8	530
2	Functional Profiling of a Plasmodium Genome Reveals an Abundance of Essential Genes. Cell, 2017, 170, 260-272.e8.	28.9	471
3	Calcium and a Calcium-Dependent Protein Kinase Regulate Gamete Formation and Mosquito Transmission in a Malaria Parasite. Cell, 2004, 117, 503-514.	28.9	415
4	A cascade of DNA-binding proteins for sexual commitment and development in Plasmodium. Nature, 2014, 507, 253-257.	27.8	366
5	Calcium-Dependent Signaling and Kinases in Apicomplexan Parasites. Cell Host and Microbe, 2009, 5, 612-622.	11.0	295
6	The conserved plant sterility gene <i>HAP2</i> functions after attachment of fusogenic membranes in <i>Chlamydomonas</i> and <i>Plasmodium</i> gametes. Genes and Development, 2008, 22, 1051-1068.	5.9	286
7	The Systematic Functional Analysis of Plasmodium Protein Kinases Identifies Essential Regulators of Mosquito Transmission. Cell Host and Microbe, 2010, 8, 377-387.	11.0	267
8	Single-cell RNA-seq and computational analysis using temporal mixture modeling resolves T _H 1/T _{FH} fate bifurcation in malaria. Science Immunology, 2017, 2, .	11.9	258
9	A comprehensive evaluation of rodent malaria parasite genomes and gene expression. BMC Biology, 2014, 12, 86.	3.8	251
10	Heparan Sulfate Proteoglycans Provide a Signal to Plasmodium Sporozoites to Stop Migrating and Productively Invade Host Cells. Cell Host and Microbe, 2007, 2, 316-327.	11.0	221
11	Gametogenesis in Malaria Parasites Is Mediated by the cGMP-Dependent Protein Kinase. PLoS Biology, 2008, 6, e139.	5.6	203
12	The Malaria Cell Atlas: Single parasite transcriptomes across the complete <i>Plasmodium</i> life cycle. Science, 2019, 365, .	12.6	198
13	Phosphoinositide Metabolism Links cGMP-Dependent Protein Kinase G to Essential Ca2+ Signals at Key Decision Points in the Life Cycle of Malaria Parasites. PLoS Biology, 2014, 12, e1001806.	5.6	185
14	A Knockout Screen of ApiAP2 Genes Reveals Networks of Interacting Transcriptional Regulators Controlling the Plasmodium Life Cycle. Cell Host and Microbe, 2017, 21, 11-22.	11.0	177
15	A Cyclic GMP Signalling Module That Regulates Gliding Motility in a Malaria Parasite. PLoS Pathogens, 2009, 5, e1000599.	4.7	171
16	Single-cell RNA-seq reveals hidden transcriptional variation in malaria parasites. ELife, 2018, 7, .	6.0	171
17	A Plasmodium Calcium-Dependent Protein Kinase Controls Zygote Development and Transmission by Translationally Activating Repressed mRNAs. Cell Host and Microbe, 2012, 12, 9-19.	11.0	163
18	Nutrient sensing modulates malaria parasite virulence. Nature, 2017, 547, 213-216.	27.8	146

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19	Plasmodium berghei calcium-dependent protein kinase 3 is required for ookinete gliding motility and mosquito midgut invasion. Molecular Microbiology, 2006, 60, 1355-1363.	2.5	141
20	The dynamics of interactions between Plasmodium and the mosquito: a study of the infectivity of Plasmodium berghei and Plasmodium gallinaceum, and their transmission by Anopheles stephensi, Anopheles gambiae and Aedes aegypti. International Journal for Parasitology, 2003, 33, 933-943.	3.1	139
21	A NIMA-related Protein Kinase Is Essential for Completion of the Sexual Cycle of Malaria Parasites. Journal of Biological Chemistry, 2005, 280, 31957-31964.	3.4	138
22	An atypical mitogen-activated protein kinase controls cytokinesis and flagellar motility during male gamete formation in a malaria parasite. Molecular Microbiology, 2005, 58, 1253-1263.	2.5	127
23	Differential recognition of members of the carcinoembryonic antigen family by Afa/Dr adhesins of diffusely adhering Escherichia coli (Afa/Dr DAEC). Molecular Microbiology, 2004, 52, 963-983.	2.5	115
24	BCKDH: The Missing Link in Apicomplexan Mitochondrial Metabolism Is Required for Full Virulence of Toxoplasma gondii and Plasmodium berghei. PLoS Pathogens, 2014, 10, e1004263.	4.7	115
25	A Genome-Scale Vector Resource Enables High-Throughput Reverse Genetic Screening in a Malaria Parasite. Cell Host and Microbe, 2015, 17, 404-413.	11.0	113
26	Protein kinases of malaria parasites: an update. Trends in Parasitology, 2008, 24, 570-577.	3.3	104
27	PlasmoGEM, a database supporting a community resource for large-scale experimental genetics in malaria parasites. Nucleic Acids Research, 2015, 43, D1176-D1182.	14.5	97
28	An Essential Role for the Plasmodium Nek-2 Nima-related Protein Kinase in the Sexual Development of Malaria Parasites. Journal of Biological Chemistry, 2009, 284, 20858-20868.	3.4	94
29	A Tetracycline-Repressible Transactivator System to Study Essential Genes in Malaria Parasites. Cell Host and Microbe, 2012, 12, 824-834.	11.0	94
30	A scalable pipeline for highly effective genetic modification of a malaria parasite. Nature Methods, 2011, 8, 1078-1082.	19.0	93
31	Genome-Scale Identification of Essential Metabolic Processes for Targeting the Plasmodium Liver Stage. Cell, 2019, 179, 1112-1128.e26.	28.9	92
32	Transmission of malaria to mosquitoes blocked by bumped kinase inhibitors. Journal of Clinical Investigation, 2012, 122, 2301-2305.	8.2	90
33	Comparative genomics in <i>Chlamydomonas</i> and <i>Plasmodium</i> identifies an ancient nuclear envelope protein family essential for sexual reproduction in protists, fungi, plants, and vertebrates. Genes and Development, 2013, 27, 1198-1215.	5.9	87
34	Both mosquito-derived xanthurenic acid and a host blood-derived factor regulate gametogenesis of Plasmodium in the midgut of the mosquito. Molecular and Biochemical Parasitology, 2001, 116, 17-24.	1.1	78
35	Protein kinases as targets for antimalarial intervention: Kinomics, structure-based design, transmission-blockade, and targeting host cell enzymes. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2005, 1754, 132-150.	2.3	78
36	Complete avian malaria parasite genomes reveal features associated with lineage-specific evolution in birds and mammals. Genome Research, 2018, 28, 547-560.	5. 5	78

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37	Inducible developmental reprogramming redefines commitment to sexual development in the malaria parasite Plasmodium berghei. Nature Microbiology, 2018, 3, 1206-1213.	13.3	77
38	Global Analysis of Apicomplexan Protein Sâ€Acyl Transferases Reveals an Enzyme Essential for Invasion. Traffic, 2013, 14, 895-911.	2.7	76
39	Azadirachtin Disrupts Formation of Organised Microtubule Arrays during Microgametogenesis of Plasmodium berghei. Journal of Eukaryotic Microbiology, 2002, 49, 489-497.	1.7	75
40	Distinct mechanisms of internalization of Neisseria gonorrhoeae by members of the CEACAM receptor family involving Rac1- and Cdc42-dependent and -independent pathways. EMBO Journal, 2002, 21, 560-571.	7.8	74
41	Calcium signalling in malaria parasites. Molecular Microbiology, 2016, 100, 397-408.	2.5	71
42	The Alveolin IMC1h Is Required for Normal Ookinete and Sporozoite Motility Behaviour and Host Colonisation in Plasmodium berghei. PLoS ONE, 2012, 7, e41409.	2.5	71
43	Invasion of hepatocytes by <i>Plasmodium</i> sporozoites requires cGMPâ€dependent protein kinase and calcium dependent protein kinase 4. Molecular Microbiology, 2016, 102, 349-363.	2.5	69
44	Mosquito cellular immunity at single-cell resolution. Science, 2020, 369, 1128-1132.	12.6	68
45	Single-cell analysis of CD4+ T-cell differentiation reveals three major cell states and progressive acceleration of proliferation. Genome Biology, 2016, 17, 103.	8.8	65
46	Defining the Range of Pathogens Susceptible to Ifitm3 Restriction Using a Knockout Mouse Model. PLoS ONE, 2013, 8, e80723.	2.5	60
47	Sub-minute Phosphoregulation of Cell Cycle Systems during Plasmodium Gamete Formation. Cell Reports, 2017, 21, 2017-2029.	6.4	59
48	Efficacy of a Plasmodium vivax Malaria Vaccine Using ChAd63 and Modified Vaccinia Ankara Expressing Thrombospondin-Related Anonymous Protein as Assessed with Transgenic Plasmodium berghei Parasites. Infection and Immunity, 2014, 82, 1277-1286.	2.2	53
49	Multiple roles for Plasmodium berghei phosphoinositide-specific phospholipase C in regulating gametocyte activation and differentiation. Cellular Microbiology, 2011, 13, 955-966.	2.1	51
50	Antibody-independent mechanisms regulate the establishment of chronic Plasmodium infection. Nature Microbiology, 2017, 2, 16276.	13.3	50
51	Epistasis studies reveal redundancy among calcium-dependent protein kinases in motility and invasion of malaria parasites. Nature Communications, 2018, 9, 4248.	12.8	50
52	The structural basis of CEACAM-receptor targeting by neisserial Opa proteins. Trends in Microbiology, 2000, 8, 258-260.	7.7	49
53	Landscape of the Plasmodium Interactome Reveals Both Conserved and Species-Specific Functionality. Cell Reports, 2019, 28, 1635-1647.e5.	6.4	49
54	Alpha-v–containing integrins are host receptors for the <i>Plasmodium falciparum</i> sporozoite surface protein, TRAP. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 4477-4482.	7.1	41

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55	Nuclear Factor-κB Directs Carcinoembryonic Antigen-related Cellular Adhesion Molecule 1 Receptor Expression inNeisseria gonorrhoeae-infected Epithelial Cells. Journal of Biological Chemistry, 2002, 277, 7438-7446.	3.4	37
56	The Malarial Serine Protease SUB1 Plays an Essential Role in Parasite Liver Stage Development. PLoS Pathogens, 2013, 9, e1003811.	4.7	34
57	Quantitative assessment of DNA replication to monitor microgametogenesis in Plasmodium berghei. Molecular and Biochemical Parasitology, 2009, 168, 172-176.	1.1	30
58	Genetic and transcriptional analysis of phosphoinositide-specific phospholipase C in Plasmodium. Experimental Parasitology, 2011, 129, 75-80.	1.2	28
59	Generation of gene targeting constructs for Plasmodium berghei by a PCR-based method amenable to high throughput applications. Molecular and Biochemical Parasitology, 2006, 145, 265-268.	1.1	27
60	Recombination-Mediated Genetic Engineering of Plasmodium berghei DNA. Methods in Molecular Biology, 2012, 923, 127-138.	0.9	27
61	Analysis of erythrocyte signalling pathways during Plasmodium falciparum infection identifies targets for host-directed antimalarial intervention. Nature Communications, 2020, 11, 4015.	12.8	27
62	An enhanced toolkit for the generation of knockout and marker-free fluorescent Plasmodium chabaudi. Wellcome Open Research, 2020, 5, 71.	1.8	23
63	Cutting Edge: The Membrane Attack Complex of Complement Is Required for the Development of Murine Experimental Cerebral Malaria. Journal of Immunology, 2011, 186, 6657-6660.	0.8	22
64	Decreased Rate of Plasma Arginine Appearance in Murine Malaria May Explain Hypoargininemia in Children With Cerebral Malaria. Journal of Infectious Diseases, 2016, 214, 1840-1849.	4.0	22
65	Gametocytes and Gametes. , 0, , 191-219.		19
66	Plasmodium Infection Is Associated with Impaired Hepatic Dimethylarginine Dimethylaminohydrolase Activity and Disruption of Nitric Oxide Synthase Inhibitor/Substrate Homeostasis. PLoS Pathogens, 2015, 11, e1005119.	4.7	18
67	Plasmodium berghei:Infectivity of Mice toAnopheles stephensiMosquitoes. Experimental Parasitology, 1996, 84, 371-379.	1.2	17
68	Isonicotinic acid hydrazide: an anti-tuberculosis drug inhibits malarial transmission in the mosquito gut. Experimental Parasitology, 2004, 106, 30-36.	1.2	17
69	Palmitoyl transferases have critical roles in the development of mosquito and liver stages of <i>Plasmodium </i> . Cellular Microbiology, 2016, 18, 1625-1641.	2.1	17
70	Signal transduction pathways induced by virulence factors of Neisseria gonorrhoeae. International Journal of Medical Microbiology, 2001, 291, 307-314.	3.6	14
71	A 39-Amino-Acid C-Terminal Truncation of GDV1 Disrupts Sexual Commitment in Plasmodium falciparum. MSphere, 2021, 6, .	2.9	14
72	Enhanced Methylation Analysis by Recovery of Unsequenceable Fragments. PLoS ONE, 2016, 11, e0152322.	2.5	13

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73	A Stem Cell Strategy Identifies Glycophorin C as a Major Erythrocyte Receptor for the Rodent Malaria Parasite Plasmodium berghei. PLoS ONE, 2016, 11, e0158238.	2.5	11
74	A single-nucleotide polymorphism in a <i>Plasmodium berghei</i> ApiAP2 transcription factor alters the development of host immunity. Science Advances, 2020, 6, eaaw6957.	10.3	10
75	An enhanced toolkit for the generation of knockout and marker-free fluorescent Plasmodium chabaudi. Wellcome Open Research, 2020, 5, 71.	1.8	10
76	Testing the impact of a single nucleotide polymorphism in a Plasmodium berghei ApiAP2 transcription factor on experimental cerebral malaria in mice. Scientific Reports, 2020, 10, 13630.	3.3	9
77	Calcium and cyclic nucleotide signaling networks in Toxoplasma gondii. , 2020, , 577-605.		6
78	A Novel Chemically Differentiated Mouse Embryonic Stem Cell-Based Model to Study Liver Stages of Plasmodium berghei. Stem Cell Reports, 2020, 14, 1123-1134.	4.8	4
79	Systematic Identification of Plasmodium Falciparum Sporozoite Membrane Protein Interactions Reveals an Essential Role for the p24AComplex in Host Infection. Molecular and Cellular Proteomics, 2021, 20, 100038.	3.8	4
80	A parasite calcium switch and Achilles' heel revealed. Nature Structural and Molecular Biology, 2010, 17, 541-543.	8.2	3
81	Landscape of the <i>Plasmodium</i> Interactome. SSRN Electronic Journal, 0, , .	0.4	1
82	The role of the cGMP-dependent protein kinase in development of the malaria parasite. BMC Pharmacology, 2009, 9, S2.	0.4	0
83	Calcium Builds Strong Host-Parasite Interactions. Cell Host and Microbe, 2015, 18, 9-10.	11.0	0
84	CRISPRing the Elephant in the Room. Cell Host and Microbe, 2018, 24, 754-755.	11.0	0
85	Genome reconstructions of metabolism of Plasmodium RBC and liver stages. Current Opinion in Microbiology, 2021, 63, 259-266.	5.1	0
86	Cracking Ali Baba's code. ELife, 2017, 6, .	6.0	0