Chris J Janse

List of Publications by Year in Descending Order

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

149	10,227	52	98
papers	citations	h-index	g-index
160	12,131 ext. citations	9.5	5.47
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
149	Suppression of Plasmodium MIF-CD74 signaling protects against severe malaria. <i>FASEB Journal</i> , 2021 , 35, e21997	0.9	O
148	Messenger RNA expressing PfCSP induces functional, protective immune responses against malaria in mice. <i>Npj Vaccines</i> , 2021 , 6, 84	9.5	11
147	Plasmodium berghei sporozoites in nonreplicative vacuole are eliminated by a PI3P-mediated autophagy-independent pathway. <i>Cellular Microbiology</i> , 2021 , 23, e13271	3.9	3
146	Screening of viral-vectored P. falciparum pre-erythrocytic candidate vaccine antigens using chimeric rodent parasites. <i>PLoS ONE</i> , 2021 , 16, e0254498	3.7	0
145	Hemozoin-mediated inflammasome activation limits long-lived anti-malarial immunity. <i>Cell Reports</i> , 2021 , 36, 109586	10.6	2
144	A universal vaccine candidate against Plasmodium vivax malaria confers protective immunity against the three PvCSP alleles. <i>Scientific Reports</i> , 2021 , 11, 17928	4.9	0
143	Generation of Novel NF135 and NF54 Lines Expressing Fluorescent Reporter Proteins Under the Control of Strong and Constitutive Promoters. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020 , 10, 270	5.9	2
142	Preclinical Development and Assessment of Viral Vectors Expressing a Fusion Antigen of Plasmodium falciparum LSA1 and LSAP2 for Efficacy against Liver-Stage Malaria. <i>Infection and Immunity</i> , 2020 , 88,	3.7	5
141	A double-blind, placebo-controlled phase 1/2a trial of the genetically attenuated malaria vaccine PfSPZ-GA1. <i>Science Translational Medicine</i> , 2020 , 12,	17.5	16
140	Generation of a Genetically Modified Chimeric Parasite Expressing Circumsporozoite Protein for Malaria Vaccine Development. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020 , 10, 591046	5.9	2
139	Artemisinin exposure at the ring or trophozoite stage impacts sexual conversion differently. <i>ELife</i> , 2020 , 9,	8.9	8
138	Combinatorial Tim-3 and PD-1 activity sustains antigen-specific Th1 cell numbers during blood-stage malaria. <i>Parasite Immunology</i> , 2020 , 42, e12723	2.2	3
137	A Hetero-Multimeric Chitinase-Containing and Ookinete-Secreted Protein Complex Involved in Mosquito Midgut Invasion. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020 , 10, 615343	5.9	3
136	Transcriptome analysis of Plasmodium berghei during exo-erythrocytic development. <i>Malaria Journal</i> , 2019 , 18, 330	3.6	18
135	A tracer-based method enables tracking of malaria parasites during human skin infection. <i>Theranostics</i> , 2019 , 9, 2768-2778	12.1	6
134	A NF54 Reporter Line Expressing mCherry-Luciferase in Gametocytes, Sporozoites, and Liver-Stages. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019 , 9, 96	5.9	7
133	Genome-Scale Identification of Essential Metabolic Processes for Targeting the Plasmodium Liver Stage. <i>Cell</i> , 2019 , 179, 1112-1128.e26	56.2	41

(2017-2018)

132	Unravelling the immune signature of Plasmodium falciparum transmission-reducing immunity. <i>Nature Communications</i> , 2018 , 9, 558	17.4	52
131	Probabilistic data integration identifies reliable gametocyte-specific proteins and transcripts in malaria parasites. <i>Scientific Reports</i> , 2018 , 8, 410	4.9	26
130	Expression of full-length Plasmodium falciparum P48/45 in P. berghei blood stages: A method to express and evaluate vaccine antigens. <i>Molecular and Biochemical Parasitology</i> , 2018 , 224, 44-49	1.9	4
129	Tailoring a Plasmodium vivax Vaccine To Enhance Efficacy through a Combination of a CSP Virus-Like Particle and TRAP Viral Vectors. <i>Infection and Immunity</i> , 2018 , 86,	3.7	18
128	Neutralization of the Plasmodium-encoded MIF ortholog confers protective immunity against malaria infection. <i>Nature Communications</i> , 2018 , 9, 2714	17.4	40
127	Adenovirus-prime and baculovirus-boost heterologous immunization achieves sterile protection against malaria sporozoite challenge in a murine model. <i>Scientific Reports</i> , 2018 , 8, 3896	4.9	8
126	Chimeric Plasmodium falciparum parasites expressing Plasmodium vivax circumsporozoite protein fail to produce salivary gland sporozoites. <i>Malaria Journal</i> , 2018 , 17, 288	3.6	7
125	OX40 Stimulation Enhances Protective Immune Responses Induced After Vaccination With Attenuated Malaria Parasites. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018 , 8, 247	5.9	8
124	DOPS Adjuvant Confers Enhanced Protection against Malaria for VLP-TRAP Based Vaccines. <i>Diseases (Basel, Switzerland)</i> , 2018 , 6,	4.4	5
123	Pre-clinical evaluation of a -based whole-sporozoite malaria vaccine candidate. <i>Npj Vaccines</i> , 2018 , 3, 54	9.5	8
122	Prime and target immunization protects against liver-stage malaria in mice. <i>Science Translational Medicine</i> , 2018 , 10,	17.5	43
121	The Plasmodium falciparum male gametocyte protein P230p, a paralog of P230, is vital for ookinete formation and mosquito transmission. <i>Scientific Reports</i> , 2018 , 8, 14902	4.9	13
120	A sporozoite-based vaccination platform against human malaria. Npj Vaccines, 2018, 3, 33	9.5	16
119	Assessment of the Plasmodium falciparum Preerythrocytic Antigen UIS3 as a Potential Candidate for a Malaria Vaccine. <i>Infection and Immunity</i> , 2017 , 85,	3.7	12
118	Evaluation of Plasmodium vivax Cell-Traversal Protein for Ookinetes and Sporozoites as a Preerythrocytic P. vivax Vaccine. <i>Vaccine Journal</i> , 2017 , 24,		13
117	Rational development of a protective P. vivax vaccine evaluated with transgenic rodent parasite challenge models. <i>Scientific Reports</i> , 2017 , 7, 46482	4.9	26
116	Signatures of malaria-associated pathology revealed by high-resolution whole-blood transcriptomics in a rodent model of malaria. <i>Scientific Reports</i> , 2017 , 7, 41722	4.9	17
115	The Plasmodium falciparum Cell-Traversal Protein for Ookinetes and Sporozoites as a Candidate for Preerythrocytic and Transmission-Blocking Vaccines. <i>Infection and Immunity</i> , 2017 , 85,	3.7	38

114	The use of transgenic parasites in malaria vaccine research. Expert Review of Vaccines, 2017, 16, 1-13	5.2	15
113	Protective immunity differs between routes of administration of attenuated malaria parasites independent of parasite liver load. <i>Scientific Reports</i> , 2017 , 7, 10372	4.9	12
112	An in vitro assay to measure antibody-mediated inhibition of P. berghei sporozoite invasion against P. falciparum antigens. <i>Scientific Reports</i> , 2017 , 7, 17011	4.9	7
111	Natural Parasite Exposure Induces Protective Human Anti-Malarial Antibodies. <i>Immunity</i> , 2017 , 47, 119	7-312299).e <u>/b</u> 0
110	Pathogenic CD8 T Cells Cause Increased Levels of VEGF-A in Experimental Malaria-Associated Acute Respiratory Distress Syndrome, but Therapeutic VEGFR Inhibition Is Not Effective. <i>Frontiers in Cellular and Infection Microbiology</i> , 2017 , 7, 416	5.9	12
109	Malaria parasite LIMP protein regulates sporozoite gliding motility and infectivity in mosquito and mammalian hosts. <i>ELife</i> , 2017 , 6,	8.9	21
108	Protective Efficacy Induced by Genetically Attenuated Mid-to-Late Liver-Stage Arresting Plasmodium berghei hrp2 Parasites. <i>American Journal of Tropical Medicine and Hygiene</i> , 2016 , 95, 378-	82 ^{3.2}	5
107	Maternally supplied S-acyl-transferase is required for crystalloid organelle formation and transmission of the malaria parasite. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 7183-8	11.5	18
106	Vital and dispensable roles of Plasmodium multidrug resistance transporters during blood- and mosquito-stage development. <i>Molecular Microbiology</i> , 2016 , 101, 78-91	4.1	9
105	Integrated transcriptomic and proteomic analyses of P. falciparum gametocytes: molecular insight into sex-specific processes and translational repression. <i>Nucleic Acids Research</i> , 2016 , 44, 6087-101	20.1	143
104	Translational Control of UIS4 Protein of the Host-Parasite Interface Is Mediated by the RNA Binding Protein Puf2 in Plasmodium berghei Sporozoites. <i>PLoS ONE</i> , 2016 , 11, e0147940	3.7	12
103	The Plasmodium PHIST and RESA-Like Protein Families of Human and Rodent Malaria Parasites. <i>PLoS ONE</i> , 2016 , 11, e0152510	3.7	11
102	CD8+ T Cells Induce Fatal Brainstem Pathology during Cerebral Malaria via Luminal Antigen-Specific Engagement of Brain Vasculature. <i>PLoS Pathogens</i> , 2016 , 12, e1006022	7.6	68
101	Rapid Generation of Marker-Free P. falciparum Fluorescent Reporter Lines Using Modified CRISPR/Cas9 Constructs and Selection Protocol. <i>PLoS ONE</i> , 2016 , 11, e0168362	3.7	29
100	Variant Exported Blood-Stage Proteins Encoded by Plasmodium Multigene Families Are Expressed in Liver Stages Where They Are Exported into the Parasitophorous Vacuole. <i>PLoS Pathogens</i> , 2016 , 12, e1005917	7.6	34
99	Multidrug ATP-binding cassette transporters are essential for hepatic development of Plasmodium sporozoites. <i>Cellular Microbiology</i> , 2016 , 18, 369-83	3.9	17
98	The machinery underlying malaria parasite virulence is conserved between rodent and human malaria parasites. <i>Nature Communications</i> , 2016 , 7, 11659	17.4	39
97	Murine Model for Preclinical Studies of Var2CSA-Mediated Pathology Associated with Malaria in Pregnancy. <i>Infection and Immunity</i> , 2016 , 84, 1761-1774	3.7	7

(2013-2015)

96	Long-term live imaging reveals cytosolic immune responses of host hepatocytes against Plasmodium infection and parasite escape mechanisms. <i>Autophagy</i> , 2015 , 11, 1561-79	10.2	84
95	Comparative assessment of vaccine vectors encoding ten malaria antigens identifies two protective liver-stage candidates. <i>Scientific Reports</i> , 2015 , 5, 11820	4.9	40
94	A Plasmodium phospholipase is involved in disruption of the liver stage parasitophorous vacuole membrane. <i>PLoS Pathogens</i> , 2015 , 11, e1004760	7.6	56
93	Replication of Plasmodium in reticulocytes can occur without hemozoin formation, resulting in chloroquine resistance. <i>Journal of Experimental Medicine</i> , 2015 , 212, 893-903	16.6	46
92	Generation of Transgenic Rodent Malaria Parasites Expressing Human Malaria Parasite Proteins. <i>Methods in Molecular Biology</i> , 2015 , 1325, 257-86	1.4	21
91	The Plasmodium palmitoyl-S-acyl-transferase DHHC2 is essential for ookinete morphogenesis and malaria transmission. <i>Scientific Reports</i> , 2015 , 5, 16034	4.9	28
90	Hybridization and pre-zygotic reproductive barriers in Plasmodium. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015 , 282, 20143027	4.4	20
89	Distinct properties of the egress-related osmiophilic bodies in male and female gametocytes of the rodent malaria parasite Plasmodium berghei. <i>Cellular Microbiology</i> , 2015 , 17, 355-68	3.9	34
88	Control of disease tolerance to malaria by nitric oxide and carbon monoxide. <i>Cell Reports</i> , 2014 , 8, 126-	36 0.6	49
87	Two Plasmodium 6-Cys family-related proteins have distinct and critical roles in liver-stage development. <i>FASEB Journal</i> , 2014 , 28, 2158-70	0.9	64
86	P. berghei telomerase subunit TERT is essential for parasite survival. <i>PLoS ONE</i> , 2014 , 9, e108930	3.7	8
85	Genome-wide RIP-Chip analysis of translational repressor-bound mRNAs in the Plasmodium gametocyte. <i>Genome Biology</i> , 2014 , 15, 493	18.3	57
84	The subcellular location of ovalbumin in Plasmodium berghei blood stages influences the magnitude of T-cell responses. <i>Infection and Immunity</i> , 2014 , 82, 4654-65	3.7	14
83	A comprehensive evaluation of rodent malaria parasite genomes and gene expression. <i>BMC Biology</i> , 2014 , 12, 86	7.3	174
82	Plasmodium falciparum Rab5B is an N-terminally myristoylated Rab GTPase that is targeted to the parasite plasma and food vacuole membranes. <i>PLoS ONE</i> , 2014 , 9, e87695	3.7	27
81	A genetically attenuated malaria vaccine candidate based on P. falciparum b9/slarp gene-deficient sporozoites. <i>ELife</i> , 2014 , 3,	8.9	53
80	Hemozoin induces lung inflammation and correlates with malaria-associated acute respiratory distress syndrome. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2013 , 48, 589-600	5.7	66
79	Plasmodium liver load following parenteral sporozoite administration in rodents. <i>Vaccine</i> , 2013 , 31, 34	10 _{‡:6} £	20

78	Loss-of-function analyses defines vital and redundant functions of the Plasmodium rhomboid protease family. <i>Molecular Microbiology</i> , 2013 , 88, 318-38	4.1	34
77	Standardization in generating and reporting genetically modified rodent malaria parasites: the RMgmDB database. <i>Methods in Molecular Biology</i> , 2013 , 923, 139-50	1.4	12
76	Bioluminescence imaging of P. berghei Schizont sequestration in rodents. <i>Methods in Molecular Biology</i> , 2013 , 923, 353-68	1.4	8
75	Quantitative analysis of Plasmodium berghei liver stages by bioluminescence imaging. <i>Methods in Molecular Biology</i> , 2013 , 923, 429-43	1.4	23
74	Proteomic and genetic analyses demonstrate that Plasmodium berghei blood stages export a large and diverse repertoire of proteins. <i>Molecular and Cellular Proteomics</i> , 2013 , 12, 426-48	7.6	52
73	Why are male malaria parasites in such a rush?: Sex-specific evolution and host-parasite interactions. <i>Evolution, Medicine and Public Health</i> , 2013 , 2013, 3-13	3	7
72	Transgenic fluorescent Plasmodium cynomolgi liver stages enable live imaging and purification of Malaria hypnozoite-forms. <i>PLoS ONE</i> , 2013 , 8, e54888	3.7	40
71	Assessing the adequacy of attenuation of genetically modified malaria parasite vaccine candidates. <i>Vaccine</i> , 2012 , 30, 2662-70	4.1	56
70	Genetic engineering of attenuated malaria parasites for vaccination. <i>Current Opinion in Biotechnology</i> , 2012 , 23, 908-16	11.4	41
69	Salivary gland-specific P. berghei reporter lines enable rapid evaluation of tissue-specific sporozoite loads in mosquitoes. <i>PLoS ONE</i> , 2012 , 7, e36376	3.7	13
69 68		3·7 7·6	13 37
	sporozoite loads in mosquitoes. <i>PLoS ONE</i> , 2012 , 7, e36376 A putative homologue of CDC20/CDH1 in the malaria parasite is essential for male gamete	7.6	
68	sporozoite loads in mosquitoes. <i>PLoS ONE</i> , 2012 , 7, e36376 A putative homologue of CDC20/CDH1 in the malaria parasite is essential for male gamete development. <i>PLoS Pathogens</i> , 2012 , 8, e1002554 Reduced CD36-dependent tissue sequestration of Plasmodium-infected erythrocytes is	7.6	37
68 67	A putative homologue of CDC20/CDH1 in the malaria parasite is essential for male gamete development. <i>PLoS Pathogens</i> , 2012 , 8, e1002554 Reduced CD36-dependent tissue sequestration of Plasmodium-infected erythrocytes is detrimental to malaria parasite growth in vivo. <i>Journal of Experimental Medicine</i> , 2012 , 209, 93-107 A Plasmodium-encoded cytokine suppresses T-cell immunity during malaria. <i>Proceedings of the</i>	7.6	37 79
68 67 66	A putative homologue of CDC20/CDH1 in the malaria parasite is essential for male gamete development. <i>PLoS Pathogens</i> , 2012 , 8, e1002554 Reduced CD36-dependent tissue sequestration of Plasmodium-infected erythrocytes is detrimental to malaria parasite growth in vivo. <i>Journal of Experimental Medicine</i> , 2012 , 209, 93-107 A Plasmodium-encoded cytokine suppresses T-cell immunity during malaria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, E2117-26 Experimentally controlled downregulation of the histone chaperone FACT in Plasmodium berghei	7.6 16.6 11.5	377955
68 67 66 65	A putative homologue of CDC20/CDH1 in the malaria parasite is essential for male gamete development. <i>PLoS Pathogens</i> , 2012 , 8, e1002554 Reduced CD36-dependent tissue sequestration of Plasmodium-infected erythrocytes is detrimental to malaria parasite growth in vivo. <i>Journal of Experimental Medicine</i> , 2012 , 209, 93-107 A Plasmodium-encoded cytokine suppresses T-cell immunity during malaria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, E2117-26 Experimentally controlled downregulation of the histone chaperone FACT in Plasmodium berghei reveals that it is critical to male gamete fertility. <i>Cellular Microbiology</i> , 2011 , 13, 1956-74 The novel putative transporter NPT1 plays a critical role in early stages of Plasmodium berghei	7.6 16.6 11.5	37795530
6867666564	A putative homologue of CDC20/CDH1 in the malaria parasite is essential for male gamete development. PLoS Pathogens, 2012, 8, e1002554 Reduced CD36-dependent tissue sequestration of Plasmodium-infected erythrocytes is detrimental to malaria parasite growth in vivo. Journal of Experimental Medicine, 2012, 209, 93-107 A Plasmodium-encoded cytokine suppresses T-cell immunity during malaria. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, E2117-26 Experimentally controlled downregulation of the histone chaperone FACT in Plasmodium berghei reveals that it is critical to male gamete fertility. Cellular Microbiology, 2011, 13, 1956-74 The novel putative transporter NPT1 plays a critical role in early stages of Plasmodium berghei sexual development. Molecular Microbiology, 2011, 81, 1343-57 A genotype and phenotype database of genetically modified malaria-parasites. Trends in	7.6 16.6 11.5 3.9 4.1	3779553022

(2009-2011)

60	Development of the piggyBac transposable system for Plasmodium berghei and its application for random mutagenesis in malaria parasites. <i>BMC Genomics</i> , 2011 , 12, 155	4.5	24
59	Rodent blood-stage Plasmodium survive in dendritic cells that infect naive mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 11205-10	11.5	44
58	CCR4-associated factor 1 coordinates the expression of Plasmodium falciparum egress and invasion proteins. <i>Eukaryotic Cell</i> , 2011 , 10, 1257-63		32
57	Aminoindoles, a novel scaffold with potent activity against Plasmodium falciparum. <i>Antimicrobial Agents and Chemotherapy</i> , 2011 , 55, 2612-22	5.9	26
56	A novel Tgene insertion/marker outT(GIMO) method for transgene expression and gene complementation in rodent malaria parasites. <i>PLoS ONE</i> , 2011 , 6, e29289	3.7	78
55	Removal of heterologous sequences from Plasmodium falciparum mutants using FLPe-recombinase. <i>PLoS ONE</i> , 2010 , 5, e15121	3.7	33
54	Glutathione reductase-null malaria parasites have normal blood stage growth but arrest during development in the mosquito. <i>Journal of Biological Chemistry</i> , 2010 , 285, 27045-27056	5.4	39
53	Three members of the 6-cys protein family of Plasmodium play a role in gamete fertility. <i>PLoS Pathogens</i> , 2010 , 6, e1000853	7.6	145
52	Novel inhibitors of Plasmodium falciparum dihydroorotate dehydrogenase with anti-malarial activity in the mouse model. <i>Journal of Biological Chemistry</i> , 2010 , 285, 33054-33064	5.4	105
51	Plasmodium falciparum dynein light chain 1 interacts with actin/myosin during blood stage development. <i>Journal of Biological Chemistry</i> , 2010 , 285, 20180-91	5.4	10
50	Universal features of post-transcriptional gene regulation are critical for Plasmodium zygote development. <i>PLoS Pathogens</i> , 2010 , 6, e1000767	7.6	180
49	Sequestration and tissue accumulation of human malaria parasites: can we learn anything from rodent models of malaria?. <i>PLoS Pathogens</i> , 2010 , 6, e1001032	7.6	80
48	Plasmepsin 4-deficient Plasmodium berghei are virulence attenuated and induce protective immunity against experimental malaria. <i>American Journal of Pathology</i> , 2010 , 176, 205-17	5.8	81
47	Visualisation and quantitative analysis of the rodent malaria liver stage by real time imaging. <i>PLoS ONE</i> , 2009 , 4, e7881	3.7	179
46	Molecular genetics and comparative genomics reveal RNAi is not functional in malaria parasites. <i>Nucleic Acids Research</i> , 2009 , 37, 3788-98	20.1	148
45	A cyclic GMP signalling module that regulates gliding motility in a malaria parasite. <i>PLoS Pathogens</i> , 2009 , 5, e1000599	7.6	133
44	The glutathione biosynthetic pathway of Plasmodium is essential for mosquito transmission. <i>PLoS Pathogens</i> , 2009 , 5, e1000302	7.6	48
43	Analysis of mutant Plasmodium berghei parasites lacking expression of multiple PbCCp genes. <i>Molecular and Biochemical Parasitology</i> , 2009 , 163, 1-7	1.9	34

42	Localisation and timing of expression of putative Plasmodium berghei rhoptry proteins in merozoites and sporozoites. <i>Molecular and Biochemical Parasitology</i> , 2009 , 166, 22-31	1.9	30
41	Egress of Plasmodium berghei gametes from their host erythrocyte is mediated by the MDV-1/PEG3 protein. <i>Cellular Microbiology</i> , 2009 , 11, 1272-88	3.9	78
40	Going live: a comparative analysis of the suitability of the RFP derivatives RedStar, mCherry and tdTomato for intravital and in vitro live imaging of Plasmodium parasites. <i>Biotechnology Journal</i> , 2009 , 4, 895-902	5.6	50
39	The Plasmodium TRAP/MIC2 family member, TRAP-Like Protein (TLP), is involved in tissue traversal by sporozoites. <i>Cellular Microbiology</i> , 2008 , 10, 1505-16	3.9	87
38	The malaria secretome: from algorithms to essential function in blood stage infection. <i>PLoS Pathogens</i> , 2008 , 4, e1000084	7.6	112
37	Proteomic profiling of Plasmodium sporozoite maturation identifies new proteins essential for parasite development and infectivity. <i>PLoS Pathogens</i> , 2008 , 4, e1000195	7.6	164
36	A conserved U-rich RNA region implicated in regulation of translation in Plasmodium female gametocytes. <i>Nucleic Acids Research</i> , 2008 , 36, 1176-86	20.1	47
35	Plasmodium lipid rafts contain proteins implicated in vesicular trafficking and signalling as well as members of the PIR superfamily, potentially implicated in host immune system interactions. <i>Proteomics</i> , 2008 , 8, 2500-13	4.8	32
34	Gene disruption of Plasmodium falciparum p52 results in attenuation of malaria liver stage development in cultured primary human hepatocytes. <i>PLoS ONE</i> , 2008 , 3, e3549	3.7	78
33	A role for natural regulatory T cells in the pathogenesis of experimental cerebral malaria. <i>American Journal of Pathology</i> , 2007 , 171, 548-59	5.8	142
32	Genetically attenuated P36p-deficient Plasmodium berghei sporozoites confer long-lasting and partial cross-species protection. <i>International Journal for Parasitology</i> , 2007 , 37, 1511-9	4.3	56
31	The exoneme helps malaria parasites to break out of blood cells. <i>Cell</i> , 2007 , 131, 1036-8	56.2	5
30	Functional characterization of the Plasmodium falciparum and P. berghei homologues of macrophage migration inhibitory factor. <i>Infection and Immunity</i> , 2007 , 75, 1116-28	3.7	69
29	High efficiency transfection of Plasmodium berghei facilitates novel selection procedures. <i>Molecular and Biochemical Parasitology</i> , 2006 , 145, 60-70	1.9	341
28	Pfs47, paralog of the male fertility factor Pfs48/45, is a female specific surface protein in Plasmodium falciparum. <i>Molecular and Biochemical Parasitology</i> , 2006 , 149, 216-22	1.9	81
27	Regulation of sexual development of Plasmodium by translational repression. <i>Science</i> , 2006 , 313, 667-9	33.3	333
26	High-efficiency transfection and drug selection of genetically transformed blood stages of the rodent malaria parasite Plasmodium berghei. <i>Nature Protocols</i> , 2006 , 1, 346-56	18.8	414
25	Real-time in vivo imaging of transgenic bioluminescent blood stages of rodent malaria parasites in mice. <i>Nature Protocols</i> , 2006 , 1, 476-85	18.8	76

(1999-2006)

24	Selection by flow-sorting of genetically transformed, GFP-expressing blood stages of the rodent malaria parasite, Plasmodium berghei. <i>Nature Protocols</i> , 2006 , 1, 614-23	18.8	81
23	Plasmodium post-genomics: better the bug you know?. <i>Nature Reviews Microbiology</i> , 2006 , 4, 344-57	22.2	58
22	Towards systematic identification of Plasmodium essential genes by transposon shuttle mutagenesis. <i>Nucleic Acids Research</i> , 2005 , 33, e174	20.1	19
21	Proteome analysis of separated male and female gametocytes reveals novel sex-specific Plasmodium biology. <i>Cell</i> , 2005 , 121, 675-87	56.2	290
20	A comprehensive survey of the Plasmodium life cycle by genomic, transcriptomic, and proteomic analyses. <i>Science</i> , 2005 , 307, 82-6	33.3	662
19	A Plasmodium whole-genome synteny map: indels and synteny breakpoints as foci for species-specific genes. <i>PLoS Pathogens</i> , 2005 , 1, e44	7.6	115
18	Murine malaria parasite sequestration: CD36 is the major receptor, but cerebral pathology is unlinked to sequestration. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 11468-73	11.5	239
17	Genetically attenuated, P36p-deficient malarial sporozoites induce protective immunity and apoptosis of infected liver cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 12194-9	11.5	217
16	A Plasmodium berghei reference line that constitutively expresses GFP at a high level throughout the complete life cycle. <i>Molecular and Biochemical Parasitology</i> , 2004 , 137, 23-33	1.9	393
15	Complement-like protein TEP1 is a determinant of vectorial capacity in the malaria vector Anopheles gambiae. <i>Cell</i> , 2004 , 116, 661-70	56.2	455
14	Malaria parasites lacking eef1a have a normal S/M phase yet grow more slowly due to a longer G1 phase. <i>Molecular Microbiology</i> , 2003 , 50, 1539-51	4.1	41
13	Genome sequence and comparative analysis of the model rodent malaria parasite Plasmodium yoelii. <i>Nature</i> , 2002 , 419, 512-9	50.4	591
12	Episomal transformation of Plasmodium berghei. <i>Methods in Molecular Medicine</i> , 2002 , 72, 305-15		3
11	Functional equivalence of structurally distinct ribosomes in the malaria parasite, Plasmodium berghei. <i>Journal of Biological Chemistry</i> , 2001 , 276, 22638-47	5.4	67
10	A central role for P48/45 in malaria parasite male gamete fertility. Cell, 2001, 104, 153-64	56.2	283
9	The selectable marker human dihydrofolate reductase enables sequential genetic manipulation of the Plasmodium berghei genome. <i>Molecular and Biochemical Parasitology</i> , 2000 , 106, 199-212	1.9	82
8	The development of genetic tools for dissecting the biology of malaria parasites. <i>Annual Review of Microbiology</i> , 2000 , 54, 157-85	17.5	87
7	Heterogeneous ribosome populations are present in Plasmodium berghei during development in its vector. <i>Molecular Microbiology</i> , 1999 , 31, 253-60	4.1	29

6	Transfection of the primate malaria parasite Plasmodium knowlesi using entirely heterologous constructs. <i>Journal of Experimental Medicine</i> , 1997 , 185, 1499-503	16.6	70
5	Circumsporozoite protein is required for development of malaria sporozoites in mosquitoes. <i>Nature</i> , 1997 , 385, 336-40	50.4	231
4	Replication, expression and segregation of plasmid-borne DNA in genetically transformed malaria parasites. <i>Molecular and Biochemical Parasitology</i> , 1997 , 86, 155-62	1.9	38
3	Computer software for testing drug susceptibility of malaria parasites. <i>Cytometry</i> , 1995 , 19, 273-81		18
2	Mechanisms of pyrimethamine resistance in two different strains of Plasmodium berghei. <i>Molecular and Biochemical Parasitology</i> , 1994 , 68, 167-71	1.9	33
1	Flow cytometric screening of blood samples for malaria parasites. <i>Cytometry</i> , 1993 , 14, 276-80		42