Claudia Andrea Daubenberger

List of Publications by Year in descending order

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128 papers 3,999 citations

36 h-index 54 g-index

139 all docs

139 docs citations

times ranked

139

6172 citing authors

#	Article	IF	Citations
1	A public antibody lineage that potently inhibits malaria infection through dual binding to the circumsporozoite protein. Nature Medicine, 2018, 24, 401-407.	15.2	183
2	A year of genomic surveillance reveals how the SARS-CoV-2 pandemic unfolded in Africa. Science, 2021, 374, 423-431.	6.0	144
3	Assessment of the novel T-cell activation marker–tuberculosis assay for diagnosis of active tuberculosis in children: a prospective proof-of-concept study. Lancet Infectious Diseases, The, 2014, 14, 931-938.	4.6	142
4	Concentration and avidity of antibodies to different circumsporozoite epitopes correlate with RTS,S/AS01E malaria vaccine efficacy. Nature Communications, 2019, 10, 2174.	5.8	123
5	Diagnostic Accuracy of Kato–Katz, FLOTAC, Baermann, and PCR Methods for the Detection of Light-Intensity Hookworm and Strongyloides stercoralis Infections in Tanzania. American Journal of Tropical Medicine and Hygiene, 2014, 90, 535-545.	0.6	119
6	Controlled Human Malaria Infection of Tanzanians by Intradermal Injection of Aseptic, Purified, Cryopreserved Plasmodium falciparum Sporozoites. American Journal of Tropical Medicine and Hygiene, 2014, 91, 471-480.	0.6	116
7	Safety, Immunogenicity, and Protective Efficacy against Controlled Human Malaria Infection of Plasmodium falciparum Sporozoite Vaccine in Tanzanian Adults. American Journal of Tropical Medicine and Hygiene, 2018, 99, 338-349.	0.6	114
8	Inactivated trivalent influenza vaccination is associated with lower mortality among patients with COVID-19 in Brazil. BMJ Evidence-Based Medicine, 2021, 26, 192-193.	1.7	107
9	Public antibodies to malaria antigens generated by two LAIR1 insertion modalities. Nature, 2017, 548, 597-601.	13.7	91
10	Transient Hyperglycemia in Patients With Tuberculosis in Tanzania: Implications for Diabetes Screening Algorithms. Journal of Infectious Diseases, 2016, 213, 1163-1172.	1.9	87
11	The adjuvant GLA-SE promotes human Tfh cell expansion and emergence of public $TCR\hat{I}^2$ clonotypes. Journal of Experimental Medicine, 2019, 216, 1857-1873.	4.2	87
12	Safety, tolerability and immunogenicity of new formulations of the Plasmodium falciparum malaria peptide vaccine SPf66 combined with the immunological adjuvant QS-21. Vaccine, 2002, 20, 2263-2277.	1.7	79
13	Sequence and diversity of DRB genes of Aotus nancymaae, a primate model for human malaria parasites. Immunogenetics, 2000, 51, 219-230.	1.2	77
14	Investigations on the interplays between Schistosoma mansoni, praziquantel and the gut microbiome. Parasites and Vectors, 2018, 11, 168.	1.0	75
15	Summary of the animal homologue section of HLDA8. Cellular Immunology, 2005, 236, 51-58.	1.4	70
16	Identification and Characterization of Heme-interacting Proteins in the Malaria Parasite, Plasmodium falciparum. Journal of Biological Chemistry, 2003, 278, 27354-27361.	1.6	68
17	The N'-Terminal Domain of Glyceraldehyde-3-Phosphate Dehydrogenase of the Apicomplexan Plasmodium falciparum Mediates GTPase Rab2-Dependent Recruitment to Membranes. Biological Chemistry, 2003, 384, 1227-37.	1.2	66
18	Baseline exposure, antibody subclass, and hepatitis B response differentially affect malaria protective immunity following RTS,S/AS01E vaccination in African children. BMC Medicine, 2018, 16, 197.	2.3	65

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#	Article	IF	CITATIONS
19	Vaccination of cattle with TickGARD induces cross-reactive antibodies binding to conserved linear peptides of Bm86 homologues in Boophilus decoloratus. Vaccine, 2007, 25, 1287-1296.	1.7	62
20	Virosome-Formulated Plasmodium falciparum AMA-1 & Derived Peptides as Malaria Vaccine: Randomized Phase 1b Trial in Semi-Immune Adults & PloS ONE, 2011, 6, e22273.	1.1	61
21	Safety and Differential Antibody and T-Cell Responses to the Plasmodium falciparum Sporozoite Malaria Vaccine, PfSPZ Vaccine, by Age in Tanzanian Adults, Adolescents, Children, and Infants. American Journal of Tropical Medicine and Hygiene, 2019, 100, 1433-1444.	0.6	61
22	Advancing Global Health through Development and Clinical Trials Partnerships: A Randomized, Placebo-Controlled, Double-Blind Assessment of Safety, Tolerability, and Immunogenicity of PfSPZ Vaccine for Malaria in Healthy Equatoguinean Men. American Journal of Tropical Medicine and Hygiene, 2018, 98, 308-318.	0.6	55
23	Immune system development varies according to age, location, and anemia in African children. Science Translational Medicine, 2020, 12, .	5.8	54
24	Safety and Immunogenicity of H1/IC31 $\hat{A}^{\text{@}}$, an Adjuvanted TB Subunit Vaccine, in HIV-Infected Adults with CD4+ Lymphocyte Counts Greater than 350 cells/mm3: A Phase II, Multi-Centre, Double-Blind, Randomized, Placebo-Controlled Trial. PLoS ONE, 2014, 9, e114602.	1.1	52
25	Sequence and diversity of MHC DQA and DQB genes of the owl monkey Aotus nancymaae. Immunogenetics, 2000, 51, 528-537.	1.2	51
26	Immunogenicity and Protective Efficacy of Radiation-Attenuated and Chemo-Attenuated PfSPZ Vaccines in Equatoguinean Adults. American Journal of Tropical Medicine and Hygiene, 2021, 104, 283-293.	0.6	49
27	Systemic suppression of interferon-Â responses in Buruli ulcer patients resolves after surgical excision of the lesions caused by the extracellular pathogen Mycobacterium ulcerans. Journal of Leukocyte Biology, 2006, 79, 1150-1156.	1.5	48
28	HIV Infection Functionally Impairs Mycobacterium tuberculosis-Specific CD4 and CD8 T-Cell Responses. Journal of Virology, 2019, 93, .	1.5	48
29	Safety, immunogenicity and efficacy of PfSPZ Vaccine against malaria in infants in western Kenya: a double-blind, randomized, placebo-controlled phase 2 trial. Nature Medicine, 2021, 27, 1636-1645.	15.2	47
30	Increase of Dose Associated With Decrease in Protection Against Controlled Human Malaria Infection by PfSPZ Vaccine in Tanzanian Adults. Clinical Infectious Diseases, 2020, 71, 2849-2857.	2.9	46
31	Bovine Î ³ δT-Cell Responses to the Intracellular Protozoan Parasite <i>Theileria parva</i> Infection and Immunity, 1999, 67, 2241-2249.	1.0	46
32	Controlled Human Malaria Infection Leads to Long-Lasting Changes in Innate and Innate-like Lymphocyte Populations. Journal of Immunology, 2017, 199, 107-118.	0.4	45
33	Immunization of Malaria-Preexposed Volunteers With PfSPZ Vaccine Elicits Long-Lived IgM Invasion-Inhibitory and Complement-Fixing Antibodies. Journal of Infectious Diseases, 2018, 217, 1569-1578.	1.9	43
34	Characterization of Invasive and Colonizing Isolates of Streptococcus agalactiae in East African Adults. Journal of Clinical Microbiology, 2011, 49, 3652-3655.	1.8	42
35	High-resolution genotyping and mapping of recombination and gene conversion in the protozoan Theileria parva using whole genome sequencing. BMC Genomics, 2012, 13, 503.	1.2	41
36	Functional and Structural Similarity of $\hat{V^{3}9}\hat{V^{2}}$ T Cells in Humans and Aotus Monkeys, a Primate Infection Model for Plasmodium falciparum Malaria. Journal of Immunology, 2001, 167, 6421-6430.	0.4	40

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37	Impact of Malaria Preexposure on Antiparasite Cellular and Humoral Immune Responses after Controlled Human Malaria Infection. Infection and Immunity, 2015, 83, 2185-2196.	1.0	40
38	Camel Streptococcus agalactiae populations are associated with specific disease complexes and acquired the tetracycline resistance gene tetM via a Tn916-like element. Veterinary Research, 2013, 44, 86.	1.1	38
39	Molecular monitoring of the diversity of human pathogenic malaria species in blood donations on Bioko Island, Equatorial Guinea. Malaria Journal, 2019, 18, 9.	0.8	35
40	Comparison of biomarker based Matrix Assisted Laser Desorption Ionization-Time of Flight Mass Spectrometry (MALDI-TOF MS) and conventional methods in the identification of clinically relevant bacteria and yeast. BMC Microbiology, 2017, 17, 128.	1.3	34
41	RTS,S/AS01E Malaria Vaccine Induces Memory and Polyfunctional T Cell Responses in a Pediatric African Phase III Trial. Frontiers in Immunology, 2017, 8, 1008.	2.2	34
42	Antigen-stimulated PBMC transcriptional protective signatures for malaria immunization. Science Translational Medicine, 2020, 12, .	5.8	33
43	The genomes of three stocks comprising the most widely utilized live sporozoite Theileria parva vaccine exhibit very different degrees and patterns of sequence divergence. BMC Genomics, 2015, 16, 729.	1.2	31
44	Sequence and diversity of T-cell receptor alpha V , J , and C genes of the owl monkey Aotus nancymaae. Immunogenetics, 1998, 48, 253-259.	1.2	29
45	Sequence and diversity of T-cell receptor \hat{l}^2 -chain V and J genes of the owl monkey Aotus nancymaae. Immunogenetics, 1999, 49, 792-799.	1.2	29
46	Identification and recombinant expression of glyceraldehyde-3-phosphate dehydrogenase of Plasmodium falciparum. Gene, 2000, 246, 255-264.	1.0	29
47	Mixed Th1 and Th2 Mycobacterium tuberculosis-specific CD4 T cell responses in patients with active pulmonary tuberculosis from Tanzania. PLoS Neglected Tropical Diseases, 2017, 11, e0005817.	1.3	29
48	Contribution of influenza immunity and virosomal-formulated synthetic peptide to cellular immune responses in a phase I subunit malaria vaccine trial. Clinical Immunology, 2008, 127, 188-197.	1.4	28
49	Matrix-assisted laser desorption/ionization time of flight mass spectrometry for comprehensive indexing of East African ixodid tick species. Parasites and Vectors, 2016, 9, 151.	1.0	28
50	Local Activation of the Innate Immune System in Buruli Ulcer Lesions. Journal of Investigative Dermatology, 2007, 127, 638-645.	0.3	27
51	The Candidate Blood-stage Malaria Vaccine P27A Induces a Robust Humoral Response in a Fast Track to the Field Phase 1 Trial in Exposed and Nonexposed Volunteers. Clinical Infectious Diseases, 2019, 68, 466-474.	2.9	27
52	Efficacy and safety of intravenous ferric carboxymaltose compared with oral iron for the treatment of iron deficiency anaemia in women after childbirth in Tanzania: a parallel-group, open-label, randomised controlled phase 3 trial. The Lancet Global Health, 2021, 9, e189-e198.	2.9	27
53	Distribution and Risk Factors for Plasmodium and Helminth Co-infections: A Cross-Sectional Survey among Children in Bagamoyo District, Coastal Region of Tanzania. PLoS Neglected Tropical Diseases, 2015, 9, e0003660.	1.3	25
54	Induction of humoral immune response to multiple recombinant Rhipicephalus appendiculatus antigens and their effect on tick feeding success and pathogen transmission. Parasites and Vectors, 2016, 9, 484.	1.0	25

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55	Distinct Helper T Cell Type 1 and 2 Responses Associated With Malaria Protection and Risk in RTS,S/AS01E Vaccinees. Clinical Infectious Diseases, 2017, 65, 746-755.	2.9	25
56	drLumi: An open-source package to manage data, calibrate, and conduct quality control of multiplex bead-based immunoassays data analysis. PLoS ONE, 2017, 12, e0187901.	1.1	25
57	Sequence and expression of MHC-DPB1 molecules of the New World monkey Aotus nancymaae, a primate model for Plasmodium falciparum. Immunogenetics, 2002, 54, 251-259.	1.2	24
58	Sequence diversity of the merozoite surface protein 1 of Plasmodium falciparum in clinical isolates from the Kilombero District, Tanzania. Acta Tropica, 2000, 74, 51-61.	0.9	21
59	Identification of a synthetic peptide inducing cross-reactive antibodies binding to Rhipicephalus (Boophilus) decoloratus, Rhipicephalus (Boophilus) microplus, Hyalomma anatolicum anatolicum and Rhipicephalus appendiculatus BM86 homologues. Vaccine, 2009, 28, 261-269.	1.7	21
60	Enterobiasis and strongyloidiasis and associated co-infections and morbidity markers in infants, preschool- and school-aged children from rural coastal Tanzania: a cross-sectional study. BMC Infectious Diseases, 2014, 14, 644.	1.3	21
61	Quantitative whole-cell MALDI-TOF MS fingerprints distinguishes human monocyte sub-populations activated by distinct microbial ligands. BMC Biotechnology, 2015, 15, 24.	1.7	19
62	A multiplex qPCR approach for detection of pfhrp2 and pfhrp3 gene deletions in multiple strain infections of Plasmodium falciparum. Scientific Reports, 2019, 9, 13107.	1.6	19
63	Whole-genome sequence-informed MALDI-TOF MS diagnostics reveal importance of Klebsiella oxytoca group in invasive infections: a retrospective clinical study. Genome Medicine, 2021, 13, 150.	3.6	19
64	Proteome-wide analysis of a malaria vaccine study reveals personalized humoral immune profiles in Tanzanian adults. ELife, 2020, 9, .	2.8	19
65	Amino acid dimorphism and parasite immune evasion: cellular immune responses to a promiscuous epitope ofPlasmodium falciparum merozoite surface protein?1 displaying dimorphic amino acid polymorphism are highly constrained. European Journal of Immunology, 2002, 32, 3667-3677.	1.6	18
66	Whole blood transcriptome changes following controlled human malaria infection in malaria pre-exposed volunteers correlate with parasite prepatent period. PLoS ONE, 2018, 13, e0199392.	1.1	18
67	Absolute Quantification of the Host-To-Parasite DNA Ratio in Theileria parva-Infected Lymphocyte Cell Lines. PLoS ONE, 2016, 11, e0150401.	1.1	17
68	Subspecies Typing of Streptococcus agalactiae Based on Ribosomal Subunit Protein Mass Variation by MALDI-TOF MS. Frontiers in Microbiology, 2019, 10, 471.	1.5	17
69	Re-annotation of the Theileria parva genome refines 53% of the proteome and uncovers essential components of N-glycosylation, a conserved pathway in many organisms. BMC Genomics, 2020, 21, 279.	1.2	17
70	Rapid Identification of SARS-CoV-2 Variants of Concern Using a Portable <i>peak</i> PCR Platform. Analytical Chemistry, 2021, 93, 16350-16359.	3.2	17
71	TLR9 agonists as adjuvants for prophylactic and therapeutic vaccines. Current Opinion in Molecular Therapeutics, 2007, 9, 45-52.	2.8	17
72	Multi-Dose Priming Regimens of PfSPZ Vaccine: Safety and Efficacy against Controlled Human Malaria Infection in Equatoguinean Adults. American Journal of Tropical Medicine and Hygiene, 2022, 106, 1215-1226.	0.6	16

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73	Generation of chimeric monoclonal antibodies from mice that carry human immunoglobulin Cl³1 heavy or Cl̂º light chain gene segments. Journal of Immunological Methods, 1998, 215, 27-37.	0.6	15
74	Functional analysis and transcriptional output of the GÃ \P ttingen minipig genome. BMC Genomics, 2015, 16, 932.	1.2	15
75	Activation of TCR VÎ 1+ and VÎ 1â 'VÎ 2â ' Î Î T Cells upon Controlled Infection with <i> Plasmodium falciparum < /i > in Tanzanian Volunteers. Journal of Immunology, 2020, 204, 180-191.</i>	0.4	14
76	Genomic Surveillance Enables the Identification of Co-infections With Multiple SARS-CoV-2 Lineages in Equatorial Guinea. Frontiers in Public Health, 2021, 9, 818401.	1.3	14
77	Proteolytic cleavage of surface proteins enhances susceptibility of lymphocytes to invasion by Theileria parva sporozoites. European Journal of Cell Biology, 1998, 76, 125-132.	1.6	13
78	The Equatoguinean Malaria Vaccine Initiative: From the Launching of a Clinical Research Platform to Malaria Elimination Planning in Central West Africa. American Journal of Tropical Medicine and Hygiene, 2020, 103, 947-954.	0.6	13
79	A rapid, single-step purification method for immunogenic members of the hsp 70 family: validation and application. Journal of Immunological Methods, 1994, 176, 255-263.	0.6	12
80	Characterization of a reduced peptide bond analogue of a promiscuous CD4 T cell epitope derived from the Plasmodium falciparum malaria vaccine candidate merozoite surface protein 1. Molecular Immunology, 2004, 41, 775-784.	1.0	12
81	Immunization of cattle with Ra86 impedes Rhipicephalus appendiculatus nymphal-to-adult molting. Ticks and Tick-borne Diseases, 2012, 3, 170-178.	1.1	12
82	Population structure and virulence gene profiles of Streptococcus agalactiae collected from different hosts worldwide. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 527-536.	1.3	11
83	Nanotechnological immunoassay for rapid label-free analysis of candidate malaria vaccines. Nanoscale, 2021, 13, 2338-2349.	2.8	11
84	Caught in action: mechanistic insights into antibody-mediated inhibition of Plasmodium merozoite invasion. Trends in Parasitology, 2009, 25, 494-497.	1.5	10
85	Antiviral Innate Immune Activation in HIV-Infected Adults Negatively Affects H1/IC31-Induced Vaccine-Specific Memory CD4 ⁺ T Cells. Vaccine Journal, 2015, 22, 688-696.	3.2	10
86	Performance of a real-time PCR approach for diagnosing Schistosoma haematobium infections of different intensity in urine samples from Zanzibar. Infectious Diseases of Poverty, 2020, 9, 128.	1.5	10
87	Theileria parasites subvert E2F signaling to stimulate leukocyte proliferation. Scientific Reports, 2020, 10, 3982.	1.6	10
88	Flow cytometric analysis on cross-reactivity of human-specific CD monoclonal antibodies with splenocytes of Aotus nancymaae, a non-human primate model for biomedical research. Veterinary Immunology and Immunopathology, 2007, 119, 14-20.	0.5	9
89	Molecular malaria surveillance using a novel protocol for extraction and analysis of nucleic acids retained on used rapid diagnostic tests. Scientific Reports, 2020, 10, 12305.	1.6	9
90	Diagnostic performance and comparison of ultrasensitive and conventional rapid diagnostic test, thick blood smear and quantitative PCR for detection of low-density Plasmodium falciparum infections during a controlled human malaria infection study in Equatorial Guinea. Malaria Journal, 2022, 21, 99.	0.8	9

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91	Molecular characterisation of a cognate 70 kDa heat shock protein of the protozoan Theileria parva. Molecular and Biochemical Parasitology, 1997, 85, 265-269.	0.5	8
92	Association between tuberculosis, diabetes and 25 hydroxyvitamin D in Tanzania: a longitudinal case control study. BMC Infectious Diseases, 2016, 16, 626.	1.3	8
93	Draft Genome Sequences of Seven Streptococcus agalactiae Strains Isolated from Camelus dromedarius at the Horn of Africa. Genome Announcements, 2017, 5, .	0.8	8
94	Structural basis of malaria RIFIN binding by LILRB1-containing antibodies. Nature, 2021, 592, 639-643.	13.7	8
95	Identification and Characterization of a Conserved, Stage-Specific Gene Product of Plasmodium falciparum Recognized by Parasite Growth Inhibitory Antibodies. Infection and Immunity, 2003, 71, 2173-2181.	1.0	7
96	Hyperglycaemia is inversely correlated with live M. bovis BCGâ€specific CD4 ⁺ T cell responses in Tanzanian adults with latent or active tuberculosis. Immunity, Inflammation and Disease, 2018, 6, 345-353.	1.3	7
97	ELIMU-MDx: a web-based, open-source platform for storage, management and analysis of diagnostic qPCR data. BioTechniques, 2020, 68, 22-27.	0.8	7
98	Analysis of nucleic acids extracted from rapid diagnostic tests reveals a significant proportion of false positive test results associated with recent malaria treatment. Malaria Journal, 2022, 21, 23.	0.8	7
99	Antigen processing and presentation by a mouse macrophage-like cell line expressing human HLA class II molecules. International Immunology, 1996, 8, 307-315.	1.8	6
100	Structural and functional characterisation of the Toll like receptor 9 of Aotus nancymaae, a non-human primate model for malaria vaccine development. Immunogenetics, 2005, 57, 283-288.	1.2	6
101	Early whole blood transcriptional responses to radiation-attenuated Plasmodium falciparum sporozoite vaccination in malaria naĀ ve and malaria pre-exposed adult volunteers. Malaria Journal, 2021, 20, 308.	0.8	6
102	Capture-based enrichment of Theileria parva DNA enables full genome assembly of first buffalo-derived strain and reveals exceptional intra-specific genetic diversity. PLoS Neglected Tropical Diseases, 2020, 14, e0008781.	1.3	6
103	Maturation and Mip- $1\hat{l}^2$ Production of Cytomegalovirus-Specific T Cell Responses in Tanzanian Children, Adolescents and Adults: Impact by HIV and Mycobacterium tuberculosis Co-Infections. PLoS ONE, 2015, 10, e0126716.	1.1	6
104	Characterising co-infections with Plasmodium spp., Mansonella perstans or Loa loa in asymptomatic children, adults and elderly people living on Bioko Island using nucleic acids extracted from malaria rapid diagnostic tests. PLoS Neglected Tropical Diseases, 2022, 16, e0009798.	1.3	6
105	Strong off-target antibody reactivity to malarial antigens induced by RTS,S/AS01E vaccination is associated with protection. JCI Insight, 2022, 7, .	2.3	6
106	Functional characterization and phenotypic monitoring of human hematopoietic stem cell expansion and differentiation of monocytes and macrophages by whole-cell mass spectrometry. Stem Cell Research, 2018, 26, 47-54.	0.3	5
107	Two cases of long-lasting, sub-microscopic Plasmodium malariae infections in adults from coastal Tanzania. Malaria Journal, 2019, 18, 149.	0.8	5
108	Red blood cell indices and prevalence of hemoglobinopathies and glucose 6 phosphate dehydrogenase deficiencies in male Tanzanian residents of Dar es Salaam. International Journal of Molecular Epidemiology and Genetics, 2014, 5, 185-94.	0.4	4

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109	Transcriptional correlates of malaria in RTS,S/ASO1-vaccinated African children: a matched case–control study. ELife, 2022, 11, .	2.8	4
110	Role of human Pegivirus infections in whole Plasmodium falciparum sporozoite vaccination and controlled human malaria infection in African volunteers. Virology Journal, 2021, 18, 28.	1.4	3
111	Incidence of Plasmodium falciparum malaria infection in 6-month to 45-year-olds on selected areas of Bioko Island, Equatorial Guinea. Malaria Journal, 2021, 20, 322.	0.8	3
112	Development of influenza virosome-based synthetic malaria vaccines. Expert Opinion on Drug Discovery, 2008, 3, 415-423.	2.5	2
113	First clinical trial of purified, irradiated malaria sporozoites in humans. Expert Review of Vaccines, 2012, 11, 31-33.	2.0	2
114	A simple, rapid typing method for Streptococcus agalactiae based on ribosomal subunit proteins by MALDI-TOF MS. Scientific Reports, 2020, 10, 8788.	1.6	2
115	Human unconventional T cells in Plasmodium falciparum infection. Seminars in Immunopathology, 2020, 42, 265-277.	2.8	2
116	Gene-expression analysis for prediction of RTS,S-induced protection in humans. Expert Review of Vaccines, 2010, 9, 465-469.	2.0	1
117	Anti-tick Vaccines for the Control of Ticks Affecting Livestock. , 2013, , 295-311.		1
118	Analysis of Nucleic Acids Extracted from Rapid Diagnostic Tests Reveals a Significant Proportion of False Positive Test Results Associated with Recent Malaria Treatment. SSRN Electronic Journal, 0, , .	0.4	1
119	Volunteer infection studies accelerate the clinical development of novel drugs against malaria. Lancet Infectious Diseases, The, 2022, 22, 753-754.	4.6	1
120	Safety and tolerance of lymph node biopsies from chronic HIV-1 volunteers in rural Tanzania. BMC Research Notes, 2019, 12, 561.	0.6	0
121	Epitope Mapping and Fine Specificity of Human T and B Cell Responses for Novel Candidate Blood-Stage Malaria Vaccine P27A. Frontiers in Immunology, 2020, 11, 412.	2.2	0
122	Towards protective immune responses against malaria in pregnant women. Lancet Infectious Diseases, The, 2020, 20, 517-519.	4.6	0
123	Assessment of experimental malaria vaccine induced protection in pre-exposed populations. Lancet Infectious Diseases, The, 2021, , .	4.6	0
124	The impact of <i>Loa loa</i> microfilaraemia on research subject retention during a whole sporozoite malaria vaccine trial in Equatorial Guinea. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2022, , .	0.7	0
125	Title is missing!. , 2020, 14, e0008781.		0
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