

Jacquin C Niles

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

2,537
citations

279798

23
h-index

315739

38
g-index

48
all docs

48
docs citations

48
times ranked

2862
citing authors

#	ARTICLE	IF	CITATIONS
1	The Plasmodium falciparum ABC transporter ABCI3 confers parasite strain-dependent pleiotropic antimalarial drug resistance. Cell Chemical Biology, 2022, 29, 824-839.e6.	5.2	14
2	Chemogenomics identifies acetyl-coenzyme A synthetase as a target for malaria treatment and prevention. Cell Chemical Biology, 2022, 29, 191-201.e8.	5.2	39
3	GeneTargeter: Automated <i>In Silico</i> Design for Genome Editing in the Malaria Parasite, <i>Plasmodium falciparum</i> . CRISPR Journal, 2022, 5, 155-164.	2.9	3
4	Functional genomics of RAP proteins and their role in mitoribosome regulation in Plasmodium falciparum. Nature Communications, 2022, 13, 1275.	12.8	12
5	Preclinical characterization and target validation of the antimalarial pantothenamide MMV693183. Nature Communications, 2022, 13, 2158.	12.8	13
6	Reaction hijacking of tyrosine tRNA synthetase as a new whole-of-life-cycle antimalarial strategy. Science, 2022, 376, 1074-1079.	12.6	25
7	Selective expression of variant surface antigens enables Plasmodium falciparum to evade immune clearance in vivo. Nature Communications, 2022, 13, .	12.8	5
8	An integrated platform for genome engineering and gene expression perturbation in Plasmodium falciparum. Scientific Reports, 2021, 11, 342.	3.3	29
9	Repositioning and Characterization of 1-(Pyridin-4-yl)pyrrolidin-2-one Derivatives as <i>Plasmodium</i> Cytoplasmic Prolyl-tRNA Synthetase Inhibitors. ACS Infectious Diseases, 2021, 7, 1680-1689.	3.8	14
10	MalDA, Accelerating Malaria Drug Discovery. Trends in Parasitology, 2021, 37, 493-507.	3.3	51
11	The antimalarial MMV688533 provides potential for single-dose cures with a high barrier to <i>Plasmodium falciparum</i> parasite resistance. Science Translational Medicine, 2021, 13, .	12.4	25
12	A newly characterized malaria antigen on erythrocyte and merozoite surfaces induces parasite inhibitory antibodies. Journal of Experimental Medicine, 2021, 218, .	8.5	2
13	Prioritization of Molecular Targets for Antimalarial Drug Discovery. ACS Infectious Diseases, 2021, 7, 2764-2776.	3.8	35
14	Targeted Covalent Inhibition of <i>Plasmodium</i> FK506 Binding Protein 35. ACS Medicinal Chemistry Letters, 2020, 11, 2131-2138.	2.8	11
15	Inhibition of Resistance-Refractory P. falciparum Kinase PKG Delivers Prophylactic, Blood Stage, and Transmission-Blocking Antiplasmodial Activity. Cell Chemical Biology, 2020, 27, 806-816.e8.	5.2	56
16	Complex nutrient channel phenotypes despite Mendelian inheritance in a Plasmodium falciparum genetic cross. PLoS Pathogens, 2020, 16, e1008363.	4.7	31
17	Assessment of Biological Role and Insight into Druggability of the <i>Plasmodium falciparum</i> Protease Plasmeprin V. ACS Infectious Diseases, 2020, 6, 738-746.	3.8	25
18	Phosphatidylinositol 3-phosphate and Hsp70 protect Plasmodium falciparum from heat-induced cell death. ELife, 2020, 9, .	6.0	20

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19	Plasmodium Niemann-Pick type C1-related protein is a druggable target required for parasite membrane homeostasis. <i>ELife</i> , 2019, 8, .	6.0	51
20	ATG8 Is Essential Specifically for an Autophagy-Independent Function in Apicoplast Biogenesis in Blood-Stage Malaria Parasites. <i>MBio</i> , 2018, 9, .	4.1	56
21	EXP2 is a nutrient-permeable channel in the vacuolar membrane of Plasmodium and is essential for protein export via PTEX. <i>Nature Microbiology</i> , 2018, 3, 1090-1098.	13.3	106
22	The chaperonin TRiC forms an oligomeric complex in the malaria parasite cytosol. <i>Cellular Microbiology</i> , 2017, 19, e12719.	2.1	56
23	Quantification of labile heme in live malaria parasites using a genetically encoded biosensor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E2068-E2076.	7.1	56
24	Plasmepsins IX and X are essential and druggable mediators of malaria parasite egress and invasion. <i>Science</i> , 2017, 358, 518-522.	12.6	152
25	Small molecule inhibition of apicomplexan FtsH1 disrupts plastid biogenesis in human pathogens. <i>ELife</i> , 2017, 6, .	6.0	47
26	Synthetic RNA-protein modules integrated with native translation mechanisms to control gene expression in malaria parasites. <i>Nature Communications</i> , 2016, 7, 10727.	12.8	157
27	A Genome-wide CRISPR Screen in Toxoplasma Identifies Essential Apicomplexan Genes. <i>Cell</i> , 2016, 166, 1423-1435.e12.	28.9	667
28	Ancient human sialic acid variant restricts an emerging zoonotic malaria parasite. <i>Nature Communications</i> , 2016, 7, 11187.	12.8	48
29	Identification of malaria parasite-infected red blood cell surface aptamers by inertial microfluidic SELEX (I-SELEX). <i>Scientific Reports</i> , 2015, 5, 11347.	3.3	57
30	Versatile control of Plasmodium falciparum gene expression with an inducible protein-RNA interaction. <i>Nature Communications</i> , 2014, 5, 5329.	12.8	44
31	Efficient CRISPR-Cas9-mediated genome editing in Plasmodium falciparum. <i>Nature Methods</i> , 2014, 11, 915-918.	19.0	205
32	An integrated strategy for efficient vector construction and multi-gene expression in Plasmodium falciparum. <i>Malaria Journal</i> , 2013, 12, 373.	2.3	18
33	Direct and specific chemical control of eukaryotic translation with a synthetic RNA-protein interaction. <i>Nucleic Acids Research</i> , 2012, 40, e64-e64.	14.5	38
34	Malarial Parasites Accumulate Labile Zinc Pools. <i>Chemistry and Biology</i> , 2012, 19, 660-661.	6.0	1
35	Inducible Control of Subcellular RNA Localization Using a Synthetic Protein-RNA Aptamer Interaction. <i>PLoS ONE</i> , 2012, 7, e46868.	2.5	6
36	Deconvolution of Microarray Data Predicts Transcriptionally Regulated Protein Kinases of Plasmodium falciparum. , 2011, , .		0

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37	Combined confocal Raman and quantitative phase microscopy system for biomedical diagnosis. Biomedical Optics Express, 2011, 2, 2484.	2.9	85
38	Peroxynitrite-induced oxidation and nitration products of guanine and 8-oxoguanine: Structures and mechanisms of product formation. Nitric Oxide - Biology and Chemistry, 2006, 14, 109-121.	2.7	173
39	Mass Spectrometric Identification of 4-Hydroxy-2,5-dioxo-imidazolidine-4-carboxylic Acid during Oxidation of 8-Oxoguanosine by Peroxynitrite and KHSO ₅ /CoCl ₂ . Chemical Research in Toxicology, 2004, 17, 1501-1509.	3.3	13
40	Spiroiminodihydantoin and Guanidinohydantoin Are the Dominant Products of 8-Oxoguanosine Oxidation at Low Fluxes of Peroxynitrite: Mechanistic Studies with ¹⁸ O. Chemical Research in Toxicology, 2004, 17, 1510-1519.	3.3	77