

Kathryn E Tiedje

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

321
citations

1307594

7
h-index

1125743

13
g-index

18
all docs

18
docs citations

18
times ranked

214
citing authors

#	ARTICLE	IF	CITATIONS
1	Evidence of strain structure in <i>Plasmodium falciparum</i> var gene repertoires in children from Gabon, West Africa. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E4103-E4111.	7.1	53
2	Competition for hosts modulates vast antigenic diversity to generate persistent strain structure in <i>Plasmodium falciparum</i> . <i>PLoS Biology</i> , 2019, 17, e3000336.	5.6	40
3	Networks of genetic similarity reveal non-neutral processes shape strain structure in <i>Plasmodium falciparum</i> . <i>Nature Communications</i> , 2018, 9, 1817.	12.8	39
4	Seasonal Variation in the Epidemiology of Asymptomatic <i>Plasmodium falciparum</i> Infections across Two Catchment Areas in Bongo District, Ghana. <i>American Journal of Tropical Medicine and Hygiene</i> , 2017, 97, 199-212.	1.4	38
5	Population genomics of virulence genes of <i>Plasmodium falciparum</i> in clinical isolates from Uganda. <i>Scientific Reports</i> , 2017, 7, 11810.	3.3	31
6	Evolutionary analyses of the major variant surface antigen-encoding genes reveal population structure of <i>Plasmodium falciparum</i> within and between continents. <i>PLoS Genetics</i> , 2021, 17, e1009269.	3.5	20
7	Evolutionary structure of <i>Plasmodium falciparum</i> major variant surface antigen genes in South America: Implications for epidemic transmission and surveillance. <i>Ecology and Evolution</i> , 2017, 7, 9376-9390.	1.9	16
8	Age-specific patterns of DBL α var diversity can explain why residents of high malaria transmission areas remain susceptible to <i>Plasmodium falciparum</i> blood stage infection throughout life. <i>International Journal for Parasitology</i> , 2022, 52, 721-731.	3.1	15
9	Frequency-Dependent Competition Between Strains Imparts Persistence to Perturbations in a Model of <i>Plasmodium falciparum</i> Malaria Transmission. <i>Frontiers in Ecology and Evolution</i> , 2021, 9, .	2.2	13
10	Lack of Geospatial Population Structure Yet Significant Linkage Disequilibrium in the Reservoir of <i>Plasmodium falciparum</i> in Bongo District, Ghana. <i>American Journal of Tropical Medicine and Hygiene</i> , 2017, 97, 1180-1189.	1.4	12
11	Indoor residual spraying with a non-pyrethroid insecticide reduces the reservoir of <i>Plasmodium falciparum</i> in a high-transmission area in northern Ghana. <i>PLOS Global Public Health</i> , 2022, 2, e0000285.	1.6	11
12	Signatures of competition and strain structure within the major blood-stage antigen of <i>Plasmodium falciparum</i> in a local community in Ghana. <i>Ecology and Evolution</i> , 2018, 8, 3574-3588.	1.9	10
13	Evolution of Antimalarial Drug Resistance Markers in the Reservoir of <i>Plasmodium falciparum</i> Infections in the Upper East Region of Ghana. <i>Journal of Infectious Diseases</i> , 2020, 222, 1692-1701.	4.0	8
14	The impact of indoor residual spraying on <i>Plasmodium falciparum</i> microsatellite variation in an area of high seasonal malaria transmission in Ghana, West Africa. <i>Molecular Ecology</i> , 2021, 30, 3974-3992.	3.9	6
15	An accurate method for identifying recent recombinants from unaligned sequences. <i>Bioinformatics</i> , 2022, 38, 1823-1829.	4.1	3