

Nationwide genetic surveillance of *Plasmodium vivax* in heterogeneous transmission dynamics and routes of migration populations

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Intra-host dynamics of co-infecting parasite genotypes in asymptomatic malaria patients. <i>Infection, Genetics and Evolution</i> , 2018, 65, 414-424.	2.3	18
2	Use cases for genetic epidemiology in malaria elimination. <i>Malaria Journal</i> , 2019, 18, 163.	2.3	57
3	Limited differentiation among <i>Plasmodium vivax</i> populations from the northwest and to the south Pacific Coast of Colombia: A malaria corridor?. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007310.	3.0	31
4	Monitoring <i>Plasmodium falciparum</i> and <i>Plasmodium vivax</i> using microsatellite markers indicates limited changes in population structure after substantial transmission decline in Papua New Guinea. <i>Molecular Ecology</i> , 2020, 29, 4525-4541.	3.9	15
5	Confirmation of the absence of local transmission and geographic assignment of imported <i>falciparum</i> malaria cases to China using microsatellite panel. <i>Malaria Journal</i> , 2020, 19, 244.	2.3	15
6	Implementing parasite genotyping into national surveillance frameworks: feedback from control programmes and researchers in the Asia-Pacific region. <i>Malaria Journal</i> , 2020, 19, 271.	2.3	31
7	SNP barcodes provide higher resolution than microsatellite markers to measure <i>Plasmodium vivax</i> population genetics. <i>Malaria Journal</i> , 2020, 19, 375.	2.3	25
8	Dynamics of <i>Plasmodium vivax</i> populations in border areas of the Greater Mekong sub-region during malaria elimination. <i>Malaria Journal</i> , 2020, 19, 145.	2.3	7
9	Genetic epidemiology of lymphatic filariasis in American Samoa after mass drug administration. <i>International Journal for Parasitology</i> , 2021, 51, 137-147.	3.1	2
10	Genomic Epidemiology in Filarial Nematodes: Transforming the Basis for Elimination Program Decisions. <i>Frontiers in Genetics</i> , 2019, 10, 1282.	2.3	29
13	Comparison of total immunoglobulin G antibody responses to different protein fragments of <i>Plasmodium vivax</i> Reticulocyte binding protein 2b. <i>Malaria Journal</i> , 2022, 21, 71.	2.3	2
14	Malaria Molecular Surveillance in the Peruvian Amazon with a Novel Highly Multiplexed <i>Plasmodium falciparum</i> AmpliSeq Assay. <i>Microbiology Spectrum</i> , 2023, 11, .	3.0	12
15	Temporal and spatial analysis of <i>Plasmodium falciparum</i> genomics reveals patterns of parasite connectivity in a low-transmission district in Southern Province, Zambia. <i>Malaria Journal</i> , 2023, 22, .	2.3	0
17	A cross-sectional study to ascertain malaria prevalence among asymptomatic travellers arriving on the Lihir Group of Islands, Papua New Guinea: implications for elimination efforts. <i>Malaria Journal</i> , 2023, 22, .	2.3	0
19	Microsatellites reveal high polymorphism and high potential for use in anti-malarial efficacy studies in areas with different transmission intensities in mainland Tanzania. <i>Malaria Journal</i> , 2024, 23, .	2.3	0