## **Guoding Zhu**

List of Publications by Year in descending order

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933447 610901 24 643 10 24 citations g-index h-index papers 24 24 24 991 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Emergence of Indigenous Artemisinin-Resistant <i>Plasmodium falciparum</i> in Africa. New England Journal of Medicine, 2017, 376, 991-993.	27.0	219
2	Relationship between Knockdown Resistance, Metabolic Detoxification and Organismal Resistance to Pyrethroids in Anopheles sinensis. PLoS ONE, 2013, 8, e55475.	2.5	61
3	Susceptibility of Anopheles sinensis to Plasmodium vivax in malarial outbreak areas of central China. Parasites and Vectors, 2013, 6, 176.	2.5	54
4	A natural symbiotic bacterium drives mosquito refractoriness to Plasmodium infection via secretion of an antimalarial lipase. Nature Microbiology, 2021, 6, 806-817.	13.3	44
5	The increasing importance of Plasmodium ovale and Plasmodium malariae in a malaria elimination setting: an observational study of imported cases in Jiangsu Province, China, 2011–2014. Malaria Journal, 2016, 15, 459.	2.3	43
6	Transcriptome profiling of pyrethroid resistant and susceptible mosquitoes in the malaria vector, Anopheles sinensis. BMC Genomics, 2014, 15, 448.	2.8	42
7	Prevalence of Drug Resistance-Associated Gene Mutations in Plasmodium vivax in Central China. Korean Journal of Parasitology, 2012, 50, 379-384.	1.3	42
8	Landscape genetic structure and evolutionary genetics of insecticide resistance gene mutations in Anopheles sinensis. Parasites and Vectors, 2016, 9, 228.	2.5	40
9	The challenge of maintaining microscopist capacity at basic levels for malaria elimination in Jiangsu Province, China. BMC Public Health, 2018, 18, 489.	2.9	25
10	Assessment of false negative rates of lactate dehydrogenase-based malaria rapid diagnostic tests for Plasmodium ovale detection. PLoS Neglected Tropical Diseases, 2019, 13, e0007254.	3.0	16
11	Limited genetic diversity of N-terminal of merozoite surface protein-1 (MSP-1) in Plasmodium ovale curtisi and P. ovale wallikeri imported from Africa to China. Parasites and Vectors, 2018, 11, 596.	2.5	8
12	Prevalence and molecular characterization of Wolbachia in field-collected Aedes albopictus, Anopheles sinensis, Armigeres subalbatus, Culex pipiens and Cx. tritaeniorhynchus in China. PLoS Neglected Tropical Diseases, 2021, 15, e0009911.	3.0	8
13	The colonization of pyrethroid resistant strain from wild Anopheles sinensis, the major Asian malaria vector. Parasites and Vectors, 2014, 7, 582.	2.5	7
14	Improving the surveillance and response system to achieve and maintain malaria elimination: a retrospective analysis in Jiangsu Province, China. Infectious Diseases of Poverty, 2022, 11, 20.	3.7	7
15	Malaria Elimination in China: Improving County-Level Malaria Personnel Knowledge of the 1-3-7 Strategy through Tabletop Exercises. American Journal of Tropical Medicine and Hygiene, 2020, 102, 804-810.	1.4	5
16	The Microbiota of Three <i>Anopheles</i> Species in China. Journal of the American Mosquito Control Association, 2021, 37, 38-40.	0.7	4
17	Differential metabolome responses to deltamethrin between resistant and susceptible Anopheles sinensis. Ecotoxicology and Environmental Safety, 2022, 237, 113553.	6.0	4
18	Plasmodium ovale curtisi and Plasmodium ovale wallikeri in Chinese travelers: Prevalence of novel genotypes of circumsporozoite protein in the African continent. Infection, Genetics and Evolution, 2019, 70, 9-14.	2.3	3

#	Article	IF	CITATION
19	Low genetic diversity and strong immunogenicity within the apical membrane antigen-1 of plasmodium ovale spp. imported from africa to china. Acta Tropica, 2020, 210, 105591.	2.0	3
20	Blood Stage of Plasmodium vivax in Central China Is Still Susceptible to Chloroquine Plus Primaquine Combination Therapy. American Journal of Tropical Medicine and Hygiene, 2013, 89, 184-187.	1.4	2
21	Immunogenicity analysis of genetically conserved segments in Plasmodium ovale merozoite surface protein-8. Parasites and Vectors, 2019, 12, 164.	2.5	2
22	A PCR-Based Technique to Track the Geographic Origin of Plasmodium falciparum With 23-SNP Barcode Analysis. Frontiers in Public Health, 2021, 9, 649170.	2.7	2
23	Genetic diversity and immunogenicity analysis of 6-cysteine protein family members in Plasmodium ovale curtisi importess from Africa to China: P12, P38 and P41. Gene Reports, 2020, 19, 100657.	0.8	1
24	Case-based malaria surveillance and response: implementation of 1-3-7 approach in Jiangsu Province, China. Advances in Parasitology, 2022, , .	3.2	1