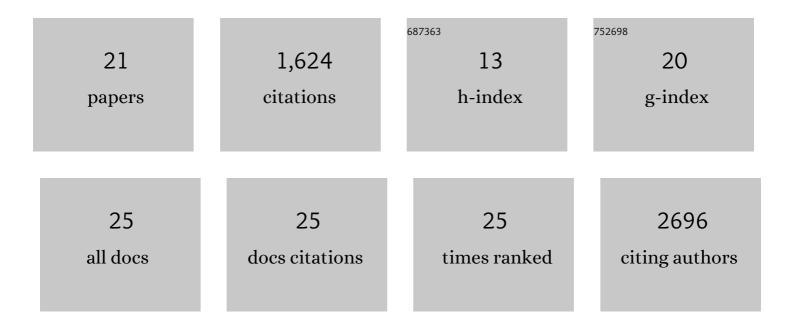
Anna E Jeffreys

List of Publications by Year in descending order

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ANNA F IFFEDEVS

#	Article	IF	CITATIONS
1	Malaria protection due to sickle haemoglobin depends on parasite genotype. Nature, 2022, 602, 106-111.	27.8	36
2	High-throughput genotyping assays for identification of glycophorin B deletion variants in population studies. Experimental Biology and Medicine, 2021, 246, 916-928.	2.4	2
3	An open dataset of Plasmodium falciparum genome variation in 7,000 worldwide samples. Wellcome Open Research, 2021, 6, 42.	1.8	97
4	An open dataset of Plasmodium falciparum genome variation in 7,000 worldwide samples. Wellcome Open Research, 2021, 6, 42.	1.8	51
5	Genetic surveillance in the Greater Mekong subregion and South Asia to support malaria control and elimination. ELife, 2021, 10, .	6.0	53
6	Haplotype heterogeneity and low linkage disequilibrium reduce reliable prediction of genotypes for the â€Î±3.7I form of α-thalassaemia using genome-wide microarray data. Wellcome Open Research, 2020, 5, 287.	1.8	3
7	Haplotype heterogeneity and low linkage disequilibrium reduce reliable prediction of genotypes for the â€î±3.7I form of î±-thalassaemia using genome-wide microarray data. Wellcome Open Research, 2020, 5, 287.	1.8	4
8	Host genetic polymorphisms and serological response against malaria in a selected population in Sri Lanka. Malaria Journal, 2018, 17, 473.	2.3	3
9	Human candidate gene polymorphisms and risk of severe malaria in children in Kilifi, Kenya: a case-control association study. Lancet Haematology,the, 2018, 5, e333-e345.	4.6	90
10	Resistance to malaria through structural variation of red blood cell invasion receptors. Science, 2017, 356, .	12.6	135
11	Malaria Host Candidate Genes Validated by Association With Current, Recent, and Historical Measures of Transmission Intensity. Journal of Infectious Diseases, 2017, 216, 45-54.	4.0	13
12	THE REAL McCOIL: A method for the concurrent estimation of the complexity of infection and SNP allele frequency for malaria parasites. PLoS Computational Biology, 2017, 13, e1005348.	3.2	93
13	Characterisation of the opposing effects of G6PD deficiency on cerebral malaria and severe malarial anaemia. ELife, 2017, 6, .	6.0	64
14	Geographic-genetic analysis of Plasmodium falciparum parasite populations from surveys of primary school children in Western Kenya. Wellcome Open Research, 2017, 2, 29.	1.8	14
15	Heterogeneous alleles comprising G6PD deficiency trait in West Africa exert contrasting effects on two major clinical presentations of severe malaria. Malaria Journal, 2016, 15, 13.	2.3	25
16	Conjunctival fibrosis and the innate barriers to Chlamydia trachomatis intracellular infection: a genome wide association study. Scientific Reports, 2015, 5, 17447.	3.3	11
17	Genetic determinants of anti-malarial acquired immunity in a large multi-centre study. Malaria Journal, 2015, 14, 333.	2.3	26
18	Imputation-Based Meta-Analysis of Severe Malaria in Three African Populations. PLoS Genetics, 2013, 9, e1003509.	3.5	95

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#	Article	IF	CITATIONS
19	Analysis of Plasmodium falciparum diversity in natural infections by deep sequencing. Nature, 2012, 487, 375-379.	27.8	450
20	Genome-wide and fine-resolution association analysis of malaria in West Africa. Nature Genetics, 2009, 41, 657-665.	21.4	345
21	Geographic-genetic analysis of Plasmodium falciparum parasite populations from surveys of primary school children in Western Kenya. Wellcome Open Research, 0, 2, 29.	1.8	10