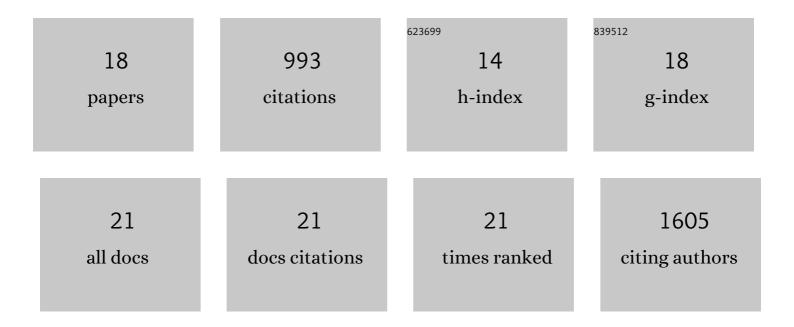
Debbie Nolder

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Distinctive genetic structure and selection patterns in Plasmodium vivax from South Asia and East Africa. Nature Communications, 2021, 12, 3160.	12.8	32
2	Failure of rapid diagnostic tests in Plasmodium falciparum malaria cases among travelers to the UK and Ireland: Identification and characterisation of the parasites. International Journal of Infectious Diseases, 2021, 108, 137-144.	3.3	12
3	Vertical Transmission of Trypanosoma Cruzi in a Non-Endemic Country: Histology of the Infected Placenta. Pediatric and Developmental Pathology, 2021, 24, 109352662110121.	1.0	1
4	Selective whole genome amplification of Plasmodium malariae DNA from clinical samples reveals insights into population structure. Scientific Reports, 2020, 10, 10832.	3.3	19
5	Chagas disease in the United Kingdom: A review of cases at the Hospital for Tropical Diseases London 1995–2018. The current state of detection of Chagas disease in the UK. Travel Medicine and Infectious Disease, 2020, 36, 101760.	3.0	9
6	A novel multiplex qPCR assay for detection of Plasmodium falciparum with histidine-rich protein 2 and 3 (pfhrp2 and pfhrp3) deletions in polyclonal infections. EBioMedicine, 2020, 55, 102757.	6.1	41
7	A molecular barcode to inform the geographical origin and transmission dynamics of Plasmodium vivax malaria. PLoS Genetics, 2020, 16, e1008576.	3.5	24
8	Geographical and temporal trends and seasonal relapse in Plasmodium ovale spp. and Plasmodium malariae infections imported to the UK between 1987 and 2015. BMC Medicine, 2018, 16, 218.	5.5	18
9	<i>pfk13</i> -Independent Treatment Failure in Four Imported Cases of Plasmodium falciparum Malaria Treated with Artemether-Lumefantrine in the United Kingdom. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	91
10	Selective Whole-Genome Amplification Is a Robust Method That Enables Scalable Whole-Genome Sequencing of <i>Plasmodium vivax</i> from Unprocessed Clinical Samples. MBio, 2017, 8, .	4.1	59
11	Paperâ€Origamiâ€Based Multiplexed Malaria Diagnostics from Whole Blood. Angewandte Chemie - International Edition, 2016, 55, 15250-15253.	13.8	125
12	Paperâ€Origamiâ€Based Multiplexed Malaria Diagnostics from Whole Blood. Angewandte Chemie, 2016, 128, 15476-15479.	2.0	29
13	Delayed Onset of Symptoms and Atovaquone-Proguanil Chemoprophylaxis Breakthrough by Plasmodium malariae in the Absence of Mutation at Codon 268 of pmcytb. PLoS Neglected Tropical Diseases, 2015, 9, e0004068.	3.0	19
14	African origin of the malaria parasite Plasmodium vivax. Nature Communications, 2014, 5, 3346.	12.8	167
15	An observational study of malaria in British travellers: <i>Plasmodium ovale wallikeri</i> and <i>Plasmodium ovale curtisi</i> differ significantly in the duration of latency. BMJ Open, 2013, 3, e002711.	1.9	69
16	Multiple hybrid genotypes of Leishmania (viannia) in a focus of mucocutaneous Leishmaniasis. American Journal of Tropical Medicine and Hygiene, 2007, 76, 573-8.	1.4	59
17	Acanthamoeba genotype T4 from the UK and Iran and isolation of the T2 genotype from clinical isolates. Journal of Medical Microbiology, 2005, 54, 755-759.	1.8	139
18	Post-mortem culture of Balamuthia mandrillaris from the brain and cerebrospinal fluid of a case of granulomatous amoebic meningoencephalitis, using human brain microvascular endothelial cells. Journal of Medical Microbiology, 2004, 53, 1007-1012.	1.8	64