Sarah Dunstan

List of Publications by Year in descending order

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83 papers 20,384 citations

76196 40 h-index 69108 77 g-index

86 all docs

86 docs citations

86 times ranked 41470 citing authors

#	Article	IF	CITATIONS
1	A global reference for human genetic variation. Nature, 2015, 526, 68-74.	13.7	13,998
2	Host Genotype-Specific Therapies Can Optimize the Inflammatory Response to Mycobacterial Infections. Cell, 2012, 148, 434-446.	13.5	523
3	The Ita4h Locus Modulates Susceptibility to Mycobacterial Infection in Zebrafish and Humans. Cell, 2010, 140, 717-730.	13.5	501
4	The Influence of Host and Bacterial Genotype on the Development of Disseminated Disease with Mycobacterium tuberculosis. PLoS Pathogens, 2008, 4, e1000034.	2.1	410
5	Genome-wide and fine-resolution association analysis of malaria in West Africa. Nature Genetics, 2009, 41, 657-665.	9.4	345
6	Reappraisal of known malaria resistance loci in a large multicenter study. Nature Genetics, 2014, 46, 1197-1204.	9.4	206
7	Human SNP Links Differential Outcomes in Inflammatory and Infectious Disease to a FOXO3-Regulated Pathway. Cell, 2013, 155, 57-69.	13.5	200
8	Salmonella: Immune Responses and Vaccines. Veterinary Journal, 2001, 161, 132-164.	0.6	190
9	Genome-wide association study identifies susceptibility loci for dengue shock syndrome at MICB and PLCE1. Nature Genetics, 2011, 43, 1139-1141.	9.4	181
10	A common human TLR1 polymorphism regulates the innate immune response to lipopeptides. European Journal of Immunology, 2007, 37, 2280-2289.	1.6	176
11	A polymorphism in human TLR2 is associated with increased susceptibility to tuberculous meningitis. Genes and Immunity, 2007, 8, 422-428.	2.2	176
12	Frequent transmission of the Mycobacterium tuberculosis Beijing lineage and positive selection for the EsxW Beijing variant in Vietnam. Nature Genetics, 2018, 50, 849-856.	9.4	167
13	A Polymorphism in Tollâ€Interleukin 1 Receptor Domain Containing Adaptor Protein Is Associated with Susceptibility to Meningeal Tuberculosis. Journal of Infectious Diseases, 2006, 194, 1127-1134.	1.9	166
14	A global network for investigating the genomic epidemiology of malaria. Nature, 2008, 456, 732-737.	13.7	148
15	Genome-wide association study identifies five new susceptibility loci for primary angle closure glaucoma. Nature Genetics, 2016, 48, 556-562.	9.4	147
16	Identification of Tuberculosis Susceptibility Genes with Human Macrophage Gene Expression Profiles. PLoS Pathogens, 2008, 4, e1000229.	2.1	134
17	Global expansion of <i>Mycobacterium tuberculosis</i> lineage 4 shaped by colonial migration and local adaptation. Science Advances, 2018, 4, eaat5869.	4.7	130
18	Molecular characterization of the staphylococcal multidrug resistance export protein QacC. Journal of Bacteriology, 1995, 177, 2827-2833.	1.0	124

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19	Copy number, linkage disequilibrium and disease association in the FCGR locus. Human Molecular Genetics, 2010, 19, 3282-3294.	1.4	119
20	Human TOLLIP Regulates TLR2 and TLR4 Signaling and Its Polymorphisms Are Associated with Susceptibility to Tuberculosis. Journal of Immunology, 2012, 189, 1737-1746.	0.4	113
21	The Role of Host Genetics in Susceptibility to Influenza: AÂSystematic Review. PLoS ONE, 2012, 7, e33180.	1.1	98
22	Genes of the Class II and Class III Major Histocompatibility Complex Are Associated with Typhoid Fever in Vietnam. Journal of Infectious Diseases, 2001, 183, 261-268.	1.9	95
23	The sensitivity of real-time PCR amplification targeting invasive Salmonellaserovars in biological specimens. BMC Infectious Diseases, 2010, 10, 125.	1.3	94
24	Leukotriene A4 Hydrolase Genotype and HIV Infection Influence Intracerebral Inflammation and Survival From Tuberculous Meningitis. Journal of Infectious Diseases, 2017, 215, 1020-1028.	1.9	93
25	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.	2.2	90
26	A Multi-Center Randomised Controlled Trial of Gatifloxacin versus Azithromycin for the Treatment of Uncomplicated Typhoid Fever in Children and Adults in Vietnam. PLoS ONE, 2008, 3, e2188.	1.1	87
27	Immune profiling with a Salmonella Typhi antigen microarray identifies new diagnostic biomarkers of human typhoid. Scientific Reports, 2013, 3, 1043.	1.6	87
28	Variation at HLA-DRB1 is associated with resistance to enteric fever. Nature Genetics, 2014, 46, 1333-1336.	9.4	85
29	The toll-like receptor 4 Asp299Gly variant: no influence on LPS responsiveness or susceptibility to pulmonary tuberculosis in The Gambia. Tuberculosis, 2004, 84, 347-352.	0.8	76
30	Transcriptional response in the peripheral blood of patients infected with Salmonella enterica serovar Typhi. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 22433-22438.	3.3	76
31	Comparison of the Abilities of Different Attenuated <i>Salmonella typhimurium</i> Strains To Elicit Humoral Immune Responses against a Heterologous Antigen. Infection and Immunity, 1998, 66, 732-740.	1.0	73
32	Characterisation of the opposing effects of G6PD deficiency on cerebral malaria and severe malarial anaemia. ELife, 2017, 6, .	2.8	64
33	Host Susceptibility and Clinical Outcomes in Tollâ€like Receptor 5–Deficient Patients with Typhoid Fever in Vietnam. Journal of Infectious Diseases, 2005, 191, 1068-1071.	1.9	61
34	The STRATAA study protocol: a programme to assess the burden of enteric fever in Bangladesh, Malawi and Nepal using prospective population census, passive surveillance, serological studies and healthcare utilisation surveys. BMJ Open, 2017, 7, e016283.	0.8	61
35	A polymorphism in human MR1 is associated with mRNA expression and susceptibility to tuberculosis. Genes and Immunity, 2017, 18, 8-14.	2.2	55
36	Use of In Vivo-Regulated Promoters To Deliver Antigens from Attenuated Salmonella enterica var. Typhimurium. Infection and Immunity, 1999, 67, 5133-5141.	1.0	52

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37	What is the evidence of a role for host genetics in susceptibility to influenza A/H5N1?. Epidemiology and Infection, 2010, 138, 1550-1558.	1.0	50
38	Common Polymorphisms in the PKP3-SIGIRR-TMEM16J Gene Region Are Associated With Susceptibility to Tuberculosis. Journal of Infectious Diseases, 2012, 205, 586-594.	1.9	50
39	Empirical ways to identify novel Bedaquiline resistance mutations in AtpE. PLoS ONE, 2019, 14, e0217169.	1.1	50
40	Human genetic variation in <i>VAC14</i> regulates <i>Salmonella</i> invasion and typhoid fever through modulation of cholesterol. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E7746-E7755.	3.3	46
41	Large scale screening for haemoglobin disorders in southern Vietnam: implications for avoidance and management. British Journal of Haematology, 2010, 150, 359-364.	1.2	44
42	TM4SF20 Ancestral Deletion and Susceptibility to a Pediatric Disorder of Early Language Delay and Cerebral White Matter Hyperintensities. American Journal of Human Genetics, 2013, 93, 197-210.	2.6	43
43	Burden of enteric fever at three urban sites in Africa and Asia: a multicentre population-based study. The Lancet Global Health, 2021, 9, e1688-e1696.	2.9	42
44	MARCO variants are associated with phagocytosis, pulmonary tuberculosis susceptibility and Beijing lineage. Genes and Immunity, 2016, 17, 419-425.	2.2	41
45	Vaccine Potential of Attenuated Mutants of Corynebacterium pseudotuberculosis in Sheep. Infection and Immunity, 1998, 66, 474-479.	1.0	40
46	DNA vaccines for bacterial infections. Immunology and Cell Biology, 1997, 75, 364-369.	1.0	39
47	Typhoid Fever and Genetic Polymorphisms at the Natural Resistance–Associated Macrophage Protein 1. Journal of Infectious Diseases, 2001, 183, 1156-1160.	1.9	39
48	Glucose-6-phosphate dehydrogenase (G6PD) mutations and haemoglobinuria syndrome in the Vietnamese population. Malaria Journal, 2009, 8, 152.	0.8	39
49	An updated systematic review of the role of host genetics in susceptibility to influenza. Influenza and Other Respiratory Viruses, 2013, 7, 37-41.	1.5	39
50	Studies of immunity and bacterial invasiveness in mice given a recombinant salmonella vector encoding murine interleukin-6. Infection and Immunity, 1996, 64, 2730-2736.	1.0	30
51	A polymorphism in human CD1A is associated with susceptibility to tuberculosis. Genes and Immunity, 2014, 15, 195-198.	2.2	29
52	In vitro and in vivo stability of recombinant plasmids in a vaccine strain of Salmonella entericavar. Typhimurium. FEMS Immunology and Medical Microbiology, 2003, 37, 111-119.	2.7	28
53	TLR9 gene region polymorphisms and susceptibility to tuberculosis in Vietnam. Tuberculosis, 2015, 95, 190-196.	0.8	27
54	Variation in human genes encoding adhesion and proinflammatory molecules are associated with severe malaria in the Vietnamese. Genes and Immunity, 2012, 13, 503-508.	2.2	24

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55	Common Polymorphisms in the CD43 Gene Region Are Associated with Tuberculosis Disease and Mortality. American Journal of Respiratory Cell and Molecular Biology, 2015, 52, 342-348.	1.4	24
56	Cytokine Release by Lipopolysaccharideâ€Stimulated Whole Blood from Patients with Typhoid Fever. Journal of Infectious Diseases, 2002, 186, 240-245.	1.9	22
57	Characterising private and shared signatures of positive selection in 37 Asian populations. European Journal of Human Genetics, 2017, 25, 499-508.	1.4	22
58	A TNF region haplotype offers protection from typhoid fever in Vietnamese patients. Human Genetics, 2007, 122, 51-61.	1.8	19
59	Second-Line HIV Therapy Outcomes and Determinants of Mortality at the Largest HIV Referral Center in Southern Vietnam. Medicine (United States), 2015, 94, e1715.	0.4	18
60	Bacterial risk factors for treatment failure and relapse among patients with isoniazid resistant tuberculosis. BMC Infectious Diseases, 2018, 18, 112.	1.3	18
61	Toll-Like Receptor 4 (TLR4) and Typhoid Fever in Vietnam. PLoS ONE, 2009, 4, e4800.	1.1	16
62	Epiregulin (EREG) variation is associated with susceptibility to tuberculosis. Genes and Immunity, 2012, 13, 275-281.	2.2	16
63	The SIGLEC14 null allele is associated with Mycobacterium tuberculosis- and BCG-induced clinical and immunologic outcomes. Tuberculosis, 2017, 104, 38-45.	0.8	16
64	High prevalence of PI resistance in patients failing second-line ART in Vietnam. Journal of Antimicrobial Chemotherapy, 2016, 71, 762-774.	1.3	15
65	Salmonella typhimurium displays normal invasion of mice with defective epidermal growth factor receptors. Infection and Immunity, 1995, 63, 2770-2772.	1.0	14
66	Whole genome–amplified DNA: insights and imputation. Nature Methods, 2008, 5, 279-280.	9.0	13
67	Sources of Multidrug Resistance in Patients With Previous Isoniazid-Resistant Tuberculosis Identified Using Whole Genome Sequencing: A Longitudinal Cohort Study. Clinical Infectious Diseases, 2020, 71, e532-e539.	2.9	13
68	Geographical distribution and disease associations of the CD45 exon 6 138G variant. Immunogenetics, 2006, 58, 235-239.	1.2	10
69	LTA4H Genotype Is Associated with Susceptibility to Bacterial Meningitis but Is Not a Critical Determinant of Outcome. PLoS ONE, 2015, 10, e0118789.	1.1	8
70	A Bayesian approach for estimating typhoid fever incidence from largeâ€scale facilityâ€based passive surveillance data. Statistics in Medicine, 2021, 40, 5853-5870.	0.8	8
71	REL and BHLHE40 Variants Are Associated with IL-12 and IL-10 Responses and Tuberculosis Risk. Journal of Immunology, 2022, 208, 1352-1361.	0.4	6
72	The immune responses to bacterial antigens encountered in vivo at mucosal surfaces. Philosophical Transactions of the Royal Society B: Biological Sciences, 2000, 355, 705-712.	1.8	5

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73	HIV-1 drug resistance in antiretroviral-naive individuals with HIV-1-associated tuberculous meningitis initiating antiretroviral therapy in Vietnam. Antiviral Therapy, 2012, 17, 905-913.	0.6	5
74	Understanding the global tuberculosis epidemic: moving towards routine whole-genome sequencing. International Journal of Tuberculosis and Lung Disease, 2019, 23, 1241-1242.	0.6	4
75	Cytokine Gene Knockout Mice—Lessons forMucosal B-Cell Development. , 1996, , 247-261.		3
76	Polymorphisms of the gene coding for copper/zinc superoxide dismutase (SOD1) in patients with Japanese encephalitis. Annals of Tropical Medicine and Parasitology, 2006, 100, 631-636.	1.6	2
77	Hiv-1 Drug Resistance in Antiretroviral-Naive Individuals with HIV-1-Associated Tuberculous Meningitis Initiating Antiretroviral Therapy in Vietnam. Antiviral Therapy, 2012, 17, 905-913.	0.6	1
78	Could omics unlock the secret of surviving tuberculous meningitis?. Lancet Infectious Diseases, The, 2018, 18, 479-480.	4.6	1
79	Studies of the pathogenesis and immunology of attenuated mutants of Salmonella enterica var. Typhimurium: lessons for human typhoid fever?. Medical Journal of Indonesia, 0, 7, 74.	0.2	0
80	Development of recombinant S-typhimurium as a model for S. typhi -based vaccine vectors. Medical Journal of Indonesia, 0, 7, 187.	0.2	0
81	Xpert Ultra and TB meningitis: advancing towards policy revision?. Annals of Infection, 2018, 2, 6-6.	0.0	0
82	Mitigating myopia in tuberculosis. Nature Immunology, 2021, 22, 675-676.	7.0	0
83	THE IMMUNE RESPONSES TO BACTERIAL ANTIGENS ENCOUNTERED <i>IN VIVO</i> AT MUCOSAL SURFACES., 2001, , .		O