Alex J Mentzer

List of Publications by Year in descending order

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57758 38395 13,591 102 44 95 citations h-index g-index papers 132 132 132 23768 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Human Cytomegalovirus and Risk of Incident Cardiovascular Disease in United Kingdom Biobank. Journal of Infectious Diseases, 2022, 225, 1179-1188.	4.0	12
2	T-cell and antibody responses to first BNT162b2 vaccine dose in previously infected and SARS-CoV-2-naive UK health-care workers: a multicentre prospective cohort study. Lancet Microbe, The, 2022, 3, e21-e31.	7.3	131
3	Serum calprotectin is not an independent predictor of severe COVID-19 in ambulatory adult patients. Journal of Infection, 2022, 84, e27-e29.	3.3	7
4	The antibody response to SARS-CoV-2 Beta underscores the antigenic distance to other variants. Cell Host and Microbe, 2022, 30, 53-68.e12.	11.0	52
5	An immunodominant NP105–113-B*07:02 cytotoxic T cell response controls viral replication and is associated with less severe COVID-19 disease. Nature Immunology, 2022, 23, 50-61.	14.5	110
6	SARS-CoV-2 Omicron-B.1.1.529 leads to widespread escape from neutralizing antibody responses. Cell, 2022, 185, 467-484.e15.	28.9	788
7	Response to Letter to the Editor by Ish et al. entitled  COVID-19 vaccine equity—the need of the hour'. QJM - Monthly Journal of the Association of Physicians, 2022, , .	0.5	O
8	Implementation and Extended Evaluation of the Euroimmun Anti-SARS-CoV-2 IgG Assay and Its Contribution to the United Kingdom's COVID-19 Public Health Response. Microbiology Spectrum, 2022, 10, e0228921.	3.0	2
9	Vitamin D Deficiency and Its Association with Iron Deficiency in African Children. Nutrients, 2022, 14, 1372.	4.1	10
10	Divergent trajectories of antiviral memory after SARS-CoV-2 infection. Nature Communications, 2022, 13, 1251.	12.8	20
11	ldentification of host–pathogen-disease relationships using a scalable multiplex serology platform in UK Biobank. Nature Communications, 2022, 13, 1818.	12.8	28
12	Why do breakthrough COVID-19 infections occur in the vaccinated?. QJM - Monthly Journal of the Association of Physicians, 2022, 115, 67-68.	0.5	2
13	Genome-Wide Association Study of Campylobacter <i>-</i> Positive Diarrhea Identifies Genes Involved in Toxin Processing and Inflammatory Response. MBio, 2022, 13, e0055622.	4.1	5
14	A loss-of-function <i>IFNAR1</i> allele in Polynesia underlies severe viral diseases in homozygotes. Journal of Experimental Medicine, 2022, 219, .	8.5	28
15	Sero-prevalence of 19 infectious pathogens and associated factors among middle-aged and elderly Chinese adults: a cross-sectional study. BMJ Open, 2022, 12, e058353.	1.9	5
16	Potent cross-reactive antibodies following Omicron breakthrough in vaccinees. Cell, 2022, 185, 2116-2131.e18.	28.9	105
17	Fatal COVID-19 outcomes are associated with an antibody response targeting epitopes shared with endemic coronaviruses. JCI Insight, 2022, 7, .	5.0	24
18	Antibody escape of SARS-CoV-2 Omicron BA.4 and BA.5 from vaccine and BA.1 serum. Cell, 2022, 185, 2422-2433.e13.	28.9	532

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19	Ancient DNA reveals five streams of migration into Micronesia and matrilocality in early Pacific seafarers. Science, 2022, 377, 72-79.	12.6	13
20	The suboptimal fibrinolytic response in COVIDâ€19 is dictated by high PAIâ€1. Journal of Thrombosis and Haemostasis, 2022, 20, 2394-2406.	3.8	30
21	Host genetics and infectious disease: new tools, insights and translational opportunities. Nature Reviews Genetics, 2021, 22, 137-153.	16.3	98
22	Phase $1/2$ trial of SARS-CoV-2 vaccine ChAdOx1 nCoV-19 with a booster dose induces multifunctional antibody responses. Nature Medicine, 2021, 27, 279-288.	30.7	265
23	T cell and antibody responses induced by a single dose of ChAdOx1 nCoV-19 (AZD1222) vaccine in a phase 1/2 clinical trial. Nature Medicine, 2021, 27, 270-278.	30.7	473
24	Malaria is a cause of iron deficiency in African children. Nature Medicine, 2021, 27, 653-658.	30.7	35
25	A haemagglutination test for rapid detection of antibodies to SARS-CoV-2. Nature Communications, 2021, 12, 1951.	12.8	54
26	T cell assays differentiate clinical and subclinical SARS-CoV-2 infections from cross-reactive antiviral responses. Nature Communications, 2021, 12, 2055.	12.8	102
27	The antigenic anatomy of SARS-CoV-2 receptor binding domain. Cell, 2021, 184, 2183-2200.e22.	28.9	331
28	Evidence of escape of SARS-CoV-2 variant B.1.351 from natural and vaccine-induced sera. Cell, 2021, 184, 2348-2361.e6.	28.9	936
29	Development and validation of the ISARIC 4C Deterioration model for adults hospitalised with COVID-19: a prospective cohort study. Lancet Respiratory Medicine, the, 2021, 9, 349-359.	10.7	161
30	Reduced neutralization of SARS-CoV-2 B.1.1.7 variant by convalescent and vaccine sera. Cell, 2021, 184, 2201-2211.e7.	28.9	442
31	Host Genome-Wide Association Study of Infant Susceptibility to <i>Shigella</i> -Associated Diarrhea. Infection and Immunity, 2021, 89, .	2.2	12
32	Antibody evasion by the P.1 strain of SARS-CoV-2. Cell, 2021, 184, 2939-2954.e9.	28.9	519
33	Prevalence and predictors of vitamin D deficiency in young African children. BMC Medicine, 2021, 19, 115.	5.5	17
34	Human genomics of the humoral immune response against polyomaviruses. Virus Evolution, 2021, 7, veab058.	4.9	9
35	Pre-existing asthma as a comorbidity does not modify cytokine responses and severity of COVID-19. Allergy, Asthma and Clinical Immunology, 2021, 17, 67.	2.0	3
36	Reduced neutralization of SARS-CoV-2 B.1.617 by vaccine and convalescent serum. Cell, 2021, 184, 4220-4236.e13.	28.9	630

3

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37	Two doses of SARS-CoV-2 vaccination induce robust immune responses to emerging SARS-CoV-2 variants of concern. Nature Communications, 2021, 12, 5061.	12.8	150
38	Discovery and validation of a three-gene signature to distinguish COVID-19 and other viral infections in emergency infectious disease presentations: a case-control and observational cohort study. Lancet Microbe, The, 2021, 2, e594-e603.	7.3	17
39	Paths and timings of the peopling of Polynesia inferred from genomic networks. Nature, 2021, 597, 522-526.	27.8	31
40	The Duration, Dynamics, and Determinants of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Antibody Responses in Individual Healthcare Workers. Clinical Infectious Diseases, 2021, 73, e699-e709.	5.8	235
41	Genetic, lifestyle, and health-related characteristics of adults without celiac disease who follow a gluten-free diet: a population-based study of 124,447 participants. American Journal of Clinical Nutrition, 2021, 113, 622-629.	4.7	12
42	Immunogenicity of standard and extended dosing intervals of BNT162b2 mRNA vaccine. Cell, 2021, 184, 5699-5714.e11.	28.9	262
43	The impact of viral mutations on recognition by SARS-CoV-2 specific TÂcells. IScience, 2021, 24, 103353.	4.1	57
44	OUP accepted manuscript. Journal of Infectious Diseases, 2021, , .	4.0	0
45	Imputation Performance in Latin American Populations: Improving Rare Variants Representation With the Inclusion of Native American Genomes. Frontiers in Genetics, 2021, 12, 719791.	2.3	7
46	Elevated risk of invasive group A streptococcal disease and host genetic variation in the human leucocyte antigen locus. Genes and Immunity, 2020, 21, 63-70.	4.1	5
47	Safety and immunogenicity of the ChAdOx1 nCoV-19 vaccine against SARS-CoV-2: a preliminary report of a phase 1/2, single-blind, randomised controlled trial. Lancet, The, 2020, 396, 467-478.	13.7	2,080
48	Characterization of human papillomavirus (HPV) 16 E6 seropositive individuals without HPV-associated malignancies after 10 years of follow-up in the UK Biobank. EBioMedicine, 2020, 62, 103123.	6.1	21
49	Distinct genetic architectures and environmental factors associate with host response to the \hat{I}^3 2-herpesvirus infections. Nature Communications, 2020, 11, 3849.	12.8	24
50	Reduced Ebola vaccine responses in CMV+ young adults is associated with expansion of CD57+KLRG1+ T cells. Journal of Experimental Medicine, 2020, 217, .	8.5	31
51	Broad and strong memory CD4+ and CD8+ T cells induced by SARS-CoV-2 in UK convalescent individuals following COVID-19. Nature Immunology, 2020, 21, 1336-1345.	14.5	1,066
52	Performance characteristics of five immunoassays for SARS-CoV-2: a head-to-head benchmark comparison. Lancet Infectious Diseases, The, 2020, 20, 1390-1400.	9.1	336
53	Genome-Wide Association Study of Cryptosporidiosis in Infants Implicates <i>PRKCA</i> . MBio, 2020, 11,	4.1	20
54	The Human Leukocyte Antigen Locus and Rheumatic Heart Disease Susceptibility in South Asians and Europeans. Scientific Reports, 2020, 10, 9004.	3.3	9

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55	Iron Deficiency Is Associated With Reduced Levels of Plasmodium falciparum-specific Antibodies in African Children. Clinical Infectious Diseases, 2020, 73, 43-49.	5.8	8
56	Estimating the burden of iron deficiency among African children. BMC Medicine, 2020, 18, 31.	5.5	47
57	Native American gene flow into Polynesia predating Easter Island settlement. Nature, 2020, 583, 572-577.	27.8	64
58	Yellow fever vaccine-associated viscerotropic disease in a 62-year-old British traveller: a case report. Journal of Travel Medicine, 2020, 27, .	3.0	3
59	Antibody testing for COVID-19: A report from theÂNational COVID Scientific Advisory Panel. Wellcome Open Research, 2020, 5, 139.	1.8	179
60	SARS-CoV-2 RNA detected in blood products from patients with COVID-19 is not associated with infectious virus. Wellcome Open Research, 2020, 5, 181.	1.8	81
61	SARS-CoV-2 RNA detected in blood products from patients with COVID-19 is not associated with infectious virus. Wellcome Open Research, 2020, 5, 181.	1.8	122
62	SARS-CoV-2 antibody prevalence, titres and neutralising activity in an antenatal cohort, United Kingdom, 14 April to 15 June 2020. Eurosurveillance, 2020, 25, .	7.0	17
63	Detection of neutralising antibodies to SARS-CoV-2 to determine population exposure in Scottish blood donors between March and May 2020. Eurosurveillance, 2020, 25, .	7.0	64
64	Differential occupational risks to healthcare workers from SARS-CoV-2 observed during a prospective observational study. ELife, 2020, 9, .	6.0	196
65	Antibodies Against <i>Chlamydia trachomatis</i> and Ovarian Cancer Risk in Two Independent Populations. Journal of the National Cancer Institute, 2019, 111, 129-136.	6.3	56
66	A genomeâ€wide association and replication study of blood pressure in Ugandan early adolescents. Molecular Genetics & Genomic Medicine, 2019, 7, e00950.	1.2	15
67	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. Cell, 2019, 179, 984-1002.e36.	28.9	152
68	The ferroportin Q248H mutation protects from anemia, but not malaria or bacteremia. Science Advances, 2019, 5, eaaw0109.	10.3	20
69	Joint sequencing of human and pathogen genomes reveals the genetics of pneumococcal meningitis. Nature Communications, 2019, 10, 2176.	12.8	83
70	HLA*LAâ€"HLA typing from linearly projected graph alignments. Bioinformatics, 2019, 35, 4394-4396.	4.1	88
71	Conservation, Extensive Heterozygosity, and Convergence of Signaling Potential All Indicate a Critical Role for KIR3DL3 in Higher Primates. Frontiers in Immunology, 2019, 10, 24.	4.8	31
72	Blood pressure risk factors in early adolescents: results from a Ugandan birth cohort. Journal of Human Hypertension, 2019, 33, 679-692.	2.2	9

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73	Iron Status and Associated Malaria Risk Among African Children. Clinical Infectious Diseases, 2019, 68, 1807-1814.	5.8	38
74	Validation of Multiplex Serology for human hepatitis viruses B and C, human T-lymphotropic virus 1 and Toxoplasma gondii. PLoS ONE, 2019, 14, e0210407.	2.5	18
75	Language continuity despite population replacement in Remote Oceania. Nature Ecology and Evolution, 2018, 2, 731-740.	7.8	91
76	Multiplex genomewide association analysis of breast milk fatty acid composition extends the phenotypic association and potential selection of <i>FADS1</i> variants to arachidonic acid, a critical infant micronutrient. Journal of Medical Genetics, 2018, 55, 459-468.	3.2	22
77	Population Turnover in Remote Oceania Shortly after Initial Settlement. Current Biology, 2018, 28, 1157-1165.e7.	3.9	91
78	Genetic variation in VAC14 is associated with bacteremia secondary to diverse pathogens in African children. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E3601-E3603.	7.1	12
79	Validation of Multiplex Serology detecting human herpesviruses 1-5. PLoS ONE, 2018, 13, e0209379.	2.5	39
80	Genome-Wide Association Study Reveals Genetic Link between Diarrhea-Associated Entamoeba histolytica Infection and Inflammatory Bowel Disease. MBio, 2018, 9, .	4.1	23
81	Long reads: their purpose and place. Human Molecular Genetics, 2018, 27, R234-R241.	2.9	249
82	The First Norovirus Longitudinal Seroepidemiological Study From Sub-Saharan Africa Reveals High Seroprevalence of Diverse Genotypes Associated With Host Susceptibility Factors. Journal of Infectious Diseases, 2018, 218, 716-725.	4.0	20
83	Th1/Th17 polarization persists following whole-cell pertussis vaccination despite repeated acellular boosters. Journal of Clinical Investigation, 2018, 128, 3853-3865.	8.2	107
84	Cohort study protocol: Bioresource in Adult Infectious Diseases (BioAID). Wellcome Open Research, 2018, 3, 97.	1.8	6
85	Association between a common immunoglobulin heavy chain allele and rheumatic heart disease risk in Oceania. Nature Communications, 2017, 8, 14946.	12.8	114
86	A Neolithic expansion, but strong genetic structure, in the independent history of New Guinea. Science, 2017, 357, 1160-1163.	12.6	45
87	The impact of prenatal exposure to parasitic infections and to anthelminthic treatment on antibody responses to routine immunisations given in infancy: Secondary analysis of a randomised controlled trial. PLoS Neglected Tropical Diseases, 2017, 11, e0005213.	3.0	18
88	A genomic history of Aboriginal Australia. Nature, 2016, 538, 207-214.	27.8	439
89	Genetic Association Analysis Reveals Differences in the Contribution of NOD2 Variants to the Clinical Phenotypes of Orofacial Granulomatosis. Inflammatory Bowel Diseases, 2016, 22, 1552-1558.	1.9	13
90	High-Accuracy HLA Type Inference from Whole-Genome Sequencing Data Using Population Reference Graphs. PLoS Computational Biology, 2016, 12, e1005151.	3.2	87

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91	Searching for the human genetic factors standing in the way of universally effective vaccines. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140341.	4.0	38
92	Optimizing the use of thiopurines in inflammatory bowel disease. Therapeutic Advances in Chronic Disease, 2015, 6, 138-146.	2.5	44
93	A mass in the liver. BMJ, The, 2013, 346, f2036-f2036.	6.0	0
94	Is it all cerebral toxoplasmosis?. Lancet, The, 2012, 379, 286.	13.7	12
95	Defects in mTR stability and telomerase activity produced by the Dkc1 A353V mutation in dyskeratosis congenita are rescued by a peptide from the dyskerin TruB domain. Clinical and Translational Oncology, 2012, 14, 755-763.	2.4	12
96	Experience with anti-TNF- $\hat{l}\pm$ therapy for orofacial granulomatosis. Journal of Oral Pathology and Medicine, 2011, 40, 14-19.	2.7	47
97	Distinguishing orofacial granulomatosis from crohn's disease: Two separate disease entities?. Inflammatory Bowel Diseases, 2011, 17, 2109-2115.	1.9	100
98	Reduced Neutralization of SARS-CoV-2 B.1.1.7 Variant from Naturally Acquired and Vaccine Induced Antibody Immunity. SSRN Electronic Journal, 0, , .	0.4	2
99	Discovery and Validation of a 3-Gene Transcriptional Signature to Distinguish COVID-19 and Other Viral Infections from Bacterial Sepsis in Adults; A Case-Control then Observational Cohort Study. SSRN Electronic Journal, 0, , .	0.4	3
100	In vivo negative regulation of SARS-CoV-2 receptor, ACE2, by interferons and its genetic control. Wellcome Open Research, 0, 6, 47.	1.8	2
101	$\hat{A}\hat{A}\hat{A}$ The Antigenic Anatomy of SARS-CoV-2 Receptor Binding Domain. SSRN Electronic Journal, 0, , .	0.4	2
102	Vitamin D Deficiency in Young African Children. SSRN Electronic Journal, 0, , .	0.4	1