

Kenneth G C Smith

List of Publications by Year in descending order

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Version: 2024-02-01

98
papers

13,129
citations

38660

50
h-index

34900

98
g-index

107
all docs

107
docs citations

107
times ranked

21548
citing authors

#	ARTICLE	IF	CITATIONS
1	B cell receptor repertoire kinetics after SARS-CoV-2 infection and vaccination. <i>Cell Reports</i> , 2022, 38, 110393.	2.9	29
2	Altered TMPRSS2 usage by SARS-CoV-2 Omicron impacts infectivity and fusogenicity. <i>Nature</i> , 2022, 603, 706-714.	13.7	756
3	Coagulation factor V is a T-cell inhibitor expressed by leukocytes in COVID-19. <i>IScience</i> , 2022, 25, 103971.	1.9	7
4	The impact of hypoxia on B cells in COVID-19. <i>EBioMedicine</i> , 2022, 77, 103878.	2.7	15
5	miR-374a-5p regulates inflammatory genes and monocyte function in patients with inflammatory bowel disease. <i>Journal of Experimental Medicine</i> , 2022, 219, .	4.2	7
6	Evidence of previous SARS-CoV-2 infection in seronegative patients with long COVID. <i>EBioMedicine</i> , 2022, 81, 104129.	2.7	21
7	A CD8+ NK cell transcriptomic signature associated with clinical outcome in relapsing remitting multiple sclerosis. <i>Nature Communications</i> , 2021, 12, 635.	5.8	33
8	SARS-CoV-2 evolution during treatment of chronic infection. <i>Nature</i> , 2021, 592, 277-282.	13.7	802
9	Homozygous <i>IL37</i> mutation associated with infantile inflammatory bowel disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	17
10	Sensitivity of SARS-CoV-2 B.1.1.7 to mRNA vaccine-elicited antibodies. <i>Nature</i> , 2021, 593, 136-141.	13.7	648
11	Transcriptional networks in at-risk individuals identify signatures of type 1 diabetes progression. <i>Science Translational Medicine</i> , 2021, 13, .	5.8	22
12	Single-cell multi-omics analysis of the immune response in COVID-19. <i>Nature Medicine</i> , 2021, 27, 904-916.	15.2	452
13	câ€Rel employs multiple mechanisms to promote the thymic development and peripheral function of regulatory T cells in mice. <i>European Journal of Immunology</i> , 2021, 51, 2006-2026.	1.6	7
14	Age-related immune response heterogeneity to SARS-CoV-2 vaccine BNT162b2. <i>Nature</i> , 2021, 596, 417-422.	13.7	549
15	Longitudinal analysis reveals that delayed bystander CD8+ T cell activation and early immune pathology distinguish severe COVID-19 from mild disease. <i>Immunity</i> , 2021, 54, 1257-1275.e8.	6.6	230
16	One Gene, Many Facets: Multiple Immune Pathway Dysregulation in SOCS1 Haploinsufficiency. <i>Frontiers in Immunology</i> , 2021, 12, 680334.	2.2	11
17	Signalling lymphocyte activation molecule family member 9 is found on select subsets of antigen-presenting cells and promotes resistance to <i>Salmonella</i> infection. <i>Immunology</i> , 2020, 159, 393-403.	2.0	7
18	Genetic feature engineering enables characterisation of shared risk factors in immune-mediated diseases. <i>Genome Medicine</i> , 2020, 12, 106.	3.6	12

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19	Dynamic regulation of hypoxia-inducible factor-1 α activity is essential for normal B cell development. <i>Nature Immunology</i> , 2020, 21, 1408-1420.	7.0	40
20	Integrative Modeling of Quantitative Plasma Lipoprotein, Metabolic, and Amino Acid Data Reveals a Multiorgan Pathological Signature of SARS-CoV-2 Infection. <i>Journal of Proteome Research</i> , 2020, 19, 4442-4454.	1.8	142
21	Treatment of COVID-19 with remdesivir in the absence of humoral immunity: a case report. <i>Nature Communications</i> , 2020, 11, 6385.	5.8	103
22	Whole-genome sequencing of a sporadic primary immunodeficiency cohort. <i>Nature</i> , 2020, 583, 90-95.	13.7	148
23	Whole-genome sequencing of patients with rare diseases in a national health system. <i>Nature</i> , 2020, 583, 96-102.	13.7	338
24	Ultrasensitive amplicon barcoding for next-generation sequencing facilitating sequence error and amplification-bias correction. <i>Scientific Reports</i> , 2020, 10, 10570.	1.6	3
25	Leupaxin Expression Is Dispensable for B Cell Immune Responses. <i>Frontiers in Immunology</i> , 2020, 11, 466.	2.2	4
26	Resolving mechanisms of immune-mediated disease in primary <sc>CD</sc> 4 T cells. <i>EMBO Molecular Medicine</i> , 2020, 12, e12112.	3.3	30
27	Screening of healthcare workers for SARS-CoV-2 highlights the role of asymptomatic carriage in COVID-19 transmission. <i>ELife</i> , 2020, 9, .	2.8	423
28	Effective control of SARS-CoV-2 transmission between healthcare workers during a period of diminished community prevalence of COVID-19. <i>ELife</i> , 2020, 9, .	2.8	40
29	The role of a functional variant of TYK2 in vasculitides and infections. <i>Clinical and Experimental Rheumatology</i> , 2020, 38, 949-955.	0.4	2
30	Genome-wide association study of eosinophilic granulomatosis with polyangiitis reveals genomic loci stratified by ANCA status. <i>Nature Communications</i> , 2019, 10, 5120.	5.8	160
31	Loss of the interleukin-6 receptor causes immunodeficiency, atopy, and abnormal inflammatory responses. <i>Journal of Experimental Medicine</i> , 2019, 216, 1986-1998.	4.2	153
32	Germline selection shapes human mitochondrial DNA diversity. <i>Science</i> , 2019, 364, .	6.0	178
33	B Cell Fc γ 3 Receptor IIb Modulates Atherosclerosis in Male and Female Mice by Controlling Adaptive Germinal Center and Innate B-1-Cell Responses. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2019, 39, 1379-1389.	1.1	17
34	Fc γ RIIb differentially regulates pre-immune and germinal center B cell tolerance in mouse and human. <i>Nature Communications</i> , 2019, 10, 1970.	5.8	20
35	Human interleukin-2 receptor β mutations associated with defects in immunity and peripheral tolerance. <i>Journal of Experimental Medicine</i> , 2019, 216, 1311-1327.	4.2	62
36	A blood-based prognostic biomarker in IBD. <i>Gut</i> , 2019, 68, 1386-1395.	6.1	132

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37	Analysis of the B cell receptor repertoire in six immune-mediated diseases. <i>Nature</i> , 2019, 574, 122-126.	13.7	178
38	EROS/CYBC1 mutations: Decreased NADPH oxidase function and chronic granulomatous disease. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, 782-785.e1.	1.5	59
39	Antibody repertoire analysis in polygenic autoimmune diseases. <i>Immunology</i> , 2018, 155, 3-17.	2.0	60
40	Metabolic exhaustion in infection, cancer and autoimmunity. <i>Nature Immunology</i> , 2018, 19, 213-221.	7.0	84
41	Cross-phenotype analysis of ImmunoChip data identifies <i>KDM4C</i> as a relevant locus for the development of systemic vasculitis. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, 589-595.	0.5	27
42	PRedicting Outcomes For Crohn's disease using a moLecular biomarkEr (PROFILE): protocol for a multicentre, randomised, biomarker-stratified trial. <i>BMJ Open</i> , 2018, 8, e026767.	0.8	55
43	Reduced monocyte and macrophage TNFSF15/TL1A expression is associated with susceptibility to inflammatory bowel disease. <i>PLoS Genetics</i> , 2018, 14, e1007458.	1.5	30
44	Combined Influence of B-Cell Receptor Rearrangement and Somatic Hypermutation on B-Cell Class-Switch Fate in Health and in Chronic Lymphocytic Leukemia. <i>Frontiers in Immunology</i> , 2018, 9, 1784.	2.2	22
45	Genome-wide association study identifies distinct genetic contributions to prognosis and susceptibility in Crohn's disease. <i>Nature Genetics</i> , 2017, 49, 262-268.	9.4	250
46	Eros is a novel transmembrane protein that controls the phagocyte respiratory burst and is essential for innate immunity. <i>Journal of Experimental Medicine</i> , 2017, 214, 1111-1128.	4.2	50
47	NBEAL2 is required for neutrophil and NK cell function and pathogen defense. <i>Journal of Clinical Investigation</i> , 2017, 127, 3521-3526.	3.9	25
48	Targeted genomic analysis reveals widespread autoimmune disease association with regulatory variants in the TNF superfamily cytokine signalling network. <i>Genome Medicine</i> , 2016, 8, 76.	3.6	17
49	T cell exhaustion and immune-mediated disease—the potential for therapeutic exhaustion. <i>Current Opinion in Immunology</i> , 2016, 43, 74-80.	2.4	66
50	T-cell exhaustion: understanding the interface of chronic viral and autoinflammatory diseases. <i>Immunology and Cell Biology</i> , 2016, 94, 935-942.	1.0	29
51	Novel aspects of autoimmunity. <i>Immunology and Cell Biology</i> , 2016, 94, 917-917.	1.0	0
52	Leucocyte subset-specific type 1 interferon signatures in SLE and other immune-mediated diseases. <i>RMD Open</i> , 2016, 2, e000183.	1.8	24
53	MT-HESS: an efficient Bayesian approach for simultaneous association detection in OMICS datasets, with application to eQTL mapping in multiple tissues. <i>Bioinformatics</i> , 2016, 32, 523-532.	1.8	25
54	T-cell exhaustion, co-stimulation and clinical outcome in autoimmunity and infection. <i>Nature</i> , 2015, 523, 612-616.	13.7	535

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55	Long-term follow-up of patients who received repeat-dose rituximab as maintenance therapy for ANCA-associated vasculitis. <i>Rheumatology</i> , 2015, 54, 1153-1160.	0.9	108
56	The Contribution of Transcriptomics to Biomarker Development in Systemic Vasculitis and SLE. <i>Current Pharmaceutical Design</i> , 2015, 21, 2225-2235.	0.9	9
57	Modules, networks and systems medicine for understanding disease and aiding diagnosis. <i>Genome Medicine</i> , 2014, 6, 82.	3.6	169
58	Prognosis in autoimmune and infectious disease: new insights from genetics. <i>Clinical and Translational Immunology</i> , 2014, 3, e15.	1.7	12
59	Fc γ RIIb inhibits immune complex-induced VEGF-A production and intranodal lymphangiogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 17971-17976.	3.3	20
60	Randomized trial of enteric-coated mycophenolate sodium versus mycophenolate mofetil in multi-system autoimmune disease. <i>CKJ: Clinical Kidney Journal</i> , 2014, 7, 562-568.	1.4	7
61	A Type I Interferon Transcriptional Signature Precedes Autoimmunity in Children Genetically at Risk for Type 1 Diabetes. <i>Diabetes</i> , 2014, 63, 2538-2550.	0.3	261
62	Human SNP Links Differential Outcomes in Inflammatory and Infectious Disease to a FOXO3-Regulated Pathway. <i>Cell</i> , 2013, 155, 57-69.	13.5	200
63	MiR-210 Is Induced by Oct-2, Regulates B Cells, and Inhibits Autoantibody Production. <i>Journal of Immunology</i> , 2013, 191, 3037-3048.	0.4	48
64	Analysis of a wild mouse promoter variant reveals a novel role for Fc γ RIIb in the control of the germinal center and autoimmunity. <i>Journal of Experimental Medicine</i> , 2012, 209, 2307-2319.	4.2	43
65	CD22 and Autoimmune Disease. <i>International Reviews of Immunology</i> , 2012, 31, 363-378.	1.5	43
66	Increased red cell turnover in a line of $\text{CD}22^{\text{deficient}}$ mice is caused by $\text{G}\pi 1^{\text{c}}$: A model for hereditary haemolytic anaemia. <i>European Journal of Immunology</i> , 2012, 42, 3212-3222.	1.6	2
67	Gene expression profiling of CD8+ T cells predicts prognosis in patients with Crohn disease and ulcerative colitis. <i>Journal of Clinical Investigation</i> , 2011, 121, 4170-4179.	3.9	268
68	Local Renal Autoantibody Production in Lupus Nephritis. <i>Journal of the American Society of Nephrology: JASN</i> , 2011, 22, 296-305.	3.0	125
69	Fc γ RIIB in autoimmunity and infection: evolutionary and therapeutic implications. <i>Nature Reviews Immunology</i> , 2010, 10, 328-343.	10.6	449
70	Fc γ RIIB, Fc γ RIIIB, and systemic lupus erythematosus. <i>Annals of the New York Academy of Sciences</i> , 2010, 1183, 69-88.	1.8	85
71	Copy number, linkage disequilibrium and disease association in the FCGR locus. <i>Human Molecular Genetics</i> , 2010, 19, 3282-3294.	1.4	119
72	A defunctioning polymorphism in FCGR2B is associated with protection against malaria but susceptibility to systemic lupus erythematosus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 7881-7885.	3.3	172

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73	Novel expression signatures identified by transcriptional analysis of separated leucocyte subsets in systemic lupus erythematosus and vasculitis. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 1208-1213.	0.5	100
74	A CD8+ T cell transcription signature predicts prognosis in autoimmune disease. <i>Nature Medicine</i> , 2010, 16, 586-591.	15.2	321
75	Low-affinity Fc γ 3 receptors, autoimmunity and infection. <i>Expert Reviews in Molecular Medicine</i> , 2009, 11, e24.	1.6	65
76	Distinct cell-specific control of autoimmunity and infection by Fc γ 3RIIb. <i>Journal of Experimental Medicine</i> , 2008, 205, 883-895.	4.2	168
77	Systemic lupus erythematosus-associated defects in the inhibitory receptor Fc γ RIIb reduce susceptibility to malaria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 7169-7174.	3.3	161
78	Fc γ RIIb controls bone marrow plasma cell persistence and apoptosis. <i>Nature Immunology</i> , 2007, 8, 419-429.	7.0	274
79	Microarray analysis of human leucocyte subsets: the advantages of positive selection and rapid purification. <i>BMC Genomics</i> , 2007, 8, 64.	1.2	93
80	Control of Rta expression critically determines transcription of viral and cellular genes following gammaherpesvirus infection. <i>Journal of General Virology</i> , 2007, 88, 1689-1697.	1.3	11
81	Long-term comparison of rituximab treatment for refractory systemic lupus erythematosus and vasculitis: Remission, relapse, and re-treatment. <i>Arthritis and Rheumatism</i> , 2006, 54, 2970-2982.	6.7	377
82	Loss of function of a lupus-associated Fc γ RIIb polymorphism through exclusion from lipid rafts. <i>Nature Medicine</i> , 2005, 11, 1056-1058.	15.2	301
83	SIGN-R1 Contributes to Protection against Lethal Pneumococcal Infection in Mice. <i>Journal of Experimental Medicine</i> , 2004, 200, 1383-1393.	4.2	144
84	B cell inhibitory receptors and autoimmunity. <i>Immunology</i> , 2003, 108, 263-273.	2.0	93
85	Interleukin 4 Reduces Expression of Inhibitory Receptors on B Cells and Abolishes CD22 and Fc γ RII-mediated B Cell Suppression. <i>Journal of Experimental Medicine</i> , 2002, 195, 1079-1085.	4.2	105
86	Growth of porcine kidneys in their native and xenograft environment. <i>Xenotransplantation</i> , 2000, 7, 96-100.	1.6	34
87	Autoimmune-prone mice share a promoter haplotype associated with reduced expression and function of the Fc receptor Fc γ RII. <i>Current Biology</i> , 2000, 10, 227-230.	1.8	222
88	Receptor Modulators of B-Cell Receptor Signalling $\hat{=}$ CD19/CD22. <i>Current Topics in Microbiology and Immunology</i> , 2000, 245, 195-212.	0.7	31
89	Apoptosis and Renal Disease. <i>Sepsis</i> , 1998, 2, 31-37.	0.5	4
90	Inhibition of the B Cell by CD22: A Requirement for Lyn. <i>Journal of Experimental Medicine</i> , 1998, 187, 807-811.	4.2	245

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91	Suppression of the humoral immune response by mycophenolate mofetil. Nephrology Dialysis Transplantation, 1998, 13, 160-164.	0.4	169
92	The extent of affinity maturation differs between the memory and antibody-forming cell compartments in the primary immune response. EMBO Journal, 1997, 16, 2996-3006.	3.5	381
93	Apoptosis and resolution of experimental renal infective tubulointerstitial nephritis. Nephrology, 1996, 2, 127-132.	0.7	14
94	CrmA expression in T lymphocytes of transgenic mice inhibits CD95 (Fas/APO-1)-transduced apoptosis, but does not cause lymphadenopathy or autoimmune disease. EMBO Journal, 1996, 15, 5167-76.	3.5	48
95	FAS is highly expressed in the germinal center but is not required for regulation of the B-cell response to antigen.. Proceedings of the National Academy of Sciences of the United States of America, 1995, 92, 11628-11632.	3.3	125
96	Soluble antigen can cause enhanced apoptosis of germinal-centre B cells. Nature, 1995, 375, 331-334.	13.7	302
97	Pregnancy-related anaemia in a haemodialysis patient treated with erythropoietin. Nephrology Dialysis Transplantation, 1993, 8, 563-564.	0.4	14
98	The potential for elderly donors to increase renal transplantation rates in Australia. Medical Journal of Australia, 1993, 158, 588-90.	0.8	1