

Jason A Greenbaum

List of Publications by Year in descending order

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Version: 2024-02-01

49
papers

12,562
citations

201385

27
h-index

243296

44
g-index

54
all docs

54
docs citations

54
times ranked

21126
citing authors

#	ARTICLE	IF	CITATIONS
1	Combined assessment of MHC binding and antigen abundance improves T cell epitope predictions. <i>IScience</i> , 2022, 25, 103850.	1.9	13
2	Single-cell eQTL analysis of activated T cell subsets reveals activation and cell type-dependent effects of disease-risk variants. <i>Science Immunology</i> , 2022, 7, eabm2508.	5.6	32
3	Towards the prediction of non-peptidic epitopes. <i>PLoS Computational Biology</i> , 2022, 18, e1009151.	1.5	2
4	Developmentally distinct CD4 ⁺ T _{reg} lineages shape the CD8 ⁺ T cell response to acute <i>Listeria</i> infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2113329119.	3.3	4
5	Targeting the tumor mutanome for personalized vaccination in a TMB low non-small cell lung cancer. , 2022, 10, e003821.		12
6	Distinguishing cell-cell complexes from dual lineage cells using single-cell transcriptomics is not trivial. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2022, 101, 547-551.	1.1	0
7	Predicting the Success of Fmoc-Based Peptide Synthesis. <i>ACS Omega</i> , 2022, 7, 23771-23781.	1.6	6
8	A comprehensive analysis of the IEDB MHC class-I automated benchmark. <i>Briefings in Bioinformatics</i> , 2022, 23, .	3.2	2
9	Comprehensive analysis of T cell immunodominance and immunoprevalence of SARS-CoV-2 epitopes in COVID-19 cases. <i>Cell Reports Medicine</i> , 2021, 2, 100204.	3.3	437
10	Transcriptome and chromatin landscape of iNKT cells are shaped by subset differentiation and antigen exposure. <i>Nature Communications</i> , 2021, 12, 1446.	5.8	21
11	TCRMatch: Predicting T-Cell Receptor Specificity Based on Sequence Similarity to Previously Characterized Receptors. <i>Frontiers in Immunology</i> , 2021, 12, 640725.	2.2	64
12	A phase 1b study of personalized neoantigen vaccine plus pembrolizumab in adults with advanced cancer.. <i>Journal of Clinical Oncology</i> , 2021, 39, 2615-2615.	0.8	4
13	Antigen-Specific Adaptive Immunity to SARS-CoV-2 in Acute COVID-19 and Associations with Age and Disease Severity. <i>Cell</i> , 2020, 183, 996-1012.e19.	13.5	1,494
14	Selective and cross-reactive SARS-CoV-2 T cell epitopes in unexposed humans. <i>Science</i> , 2020, 370, 89-94.	6.0	1,036
15	TLR9 Sensing of Self-DNA Controls Cell-Mediated Immunity to <i>Listeria</i> Infection via Rapid Conversion of Conventional CD4 ⁺ T Cells to Treg. <i>Cell Reports</i> , 2020, 31, 107249.	2.9	9
16	A behind-the-scenes tour of the IEDB curation process: an optimized process empirically integrating automation and human curation efforts. <i>Immunology</i> , 2020, 161, 139-147.	2.0	6
17	Targets of T Cell Responses to SARS-CoV-2 Coronavirus in Humans with COVID-19 Disease and Unexposed Individuals. <i>Cell</i> , 2020, 181, 1489-1501.e15.	13.5	3,220
18	Identifying inaccuracies in gene expression estimates from unstranded RNA-seq data. <i>Scientific Reports</i> , 2019, 9, 16342.	1.6	9

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19	IEDB-AR: immune epitope database analysis resource in 2019. <i>Nucleic Acids Research</i> , 2019, 47, W502-W506.	6.5	247
20	Circulating T cell-monocyte complexes are markers of immune perturbations. <i>ELife</i> , 2019, 8, .	2.8	67
21	Development of Asthma in Inner-City Children: Possible Roles of MAIT Cells and Variation in the Home Environment. <i>Journal of Immunology</i> , 2018, 200, 1995-2003.	0.4	38
22	Improved methods for predicting peptide binding affinity to MHC class II molecules. <i>Immunology</i> , 2018, 154, 394-406.	2.0	629
23	Transcriptomic Analysis of CD4+ T Cells Reveals Novel Immune Signatures of Latent Tuberculosis. <i>Journal of Immunology</i> , 2018, 200, 3283-3290.	0.4	43
24	Impact of Genetic Polymorphisms on Human Immune Cell Gene Expression. <i>Cell</i> , 2018, 175, 1701-1715.e16.	13.5	588
25	Predicting T cell recognition of MHC class I restricted neoepitopes. <i>Oncolmmunology</i> , 2018, 7, e1492508.	2.1	82
26	Response to Comment on "Development of Asthma in Inner-City Children: Possible Roles of MAIT Cells and Variation in the Home Environment". <i>Journal of Immunology</i> , 2018, 200, 3317-3318.	0.4	3
27	YAP and MRTF-A, transcriptional co-activators of RhoA-mediated gene expression, are critical for glioblastoma tumorigenicity. <i>Oncogene</i> , 2018, 37, 5492-5507.	2.6	49
28	Unique phenotypes and clonal expansions of human CD4 effector memory T cells re-expressing CD45RA. <i>Nature Communications</i> , 2017, 8, 1473.	5.8	208
29	Immunoproteomic analysis of house dust mite antigens reveals distinct classes of dominant T cell antigens according to function and serological reactivity. <i>Clinical and Experimental Allergy</i> , 2017, 47, 577-592.	1.4	26
30	T cell recognition is shaped by epitope sequence conservation in the host proteome and microbiome. <i>Immunology</i> , 2016, 148, 34-39.	2.0	31
31	Immunodominance in allergic T-cell reactivity to Japanese cedar in different geographic cohorts. <i>Annals of Allergy, Asthma and Immunology</i> , 2016, 117, 680-689.e1.	0.5	14
32	T-cell epitope conservation across allergen species is a major determinant of immunogenicity. <i>Journal of Allergy and Clinical Immunology</i> , 2016, 138, 571-578.e7.	1.5	40
33	Immunological consequences of intragenus conservation of <i>Mycobacterium tuberculosis</i> T-cell epitopes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E147-55.	3.3	69
34	Automated benchmarking of peptide-MHC class I binding predictions. <i>Bioinformatics</i> , 2015, 31, 2174-2181.	1.8	127
35	Development and validation of a sample sparing strategy for HLA typing utilizing next generation sequencing. <i>Human Immunology</i> , 2015, 76, 917-922.	1.2	11
36	IL-10-producing intestinal macrophages prevent excessive antibacterial innate immunity by limiting IL-23 synthesis. <i>Nature Communications</i> , 2015, 6, 7055.	5.8	103

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37	Dengue Virus Evolution under a Host-Targeted Antiviral. <i>Journal of Virology</i> , 2015, 89, 5592-5601.	1.5	49
38	The immune epitope database (IEDB) 3.0. <i>Nucleic Acids Research</i> , 2015, 43, D405-D412.	6.5	1,014
39	Comprehensive analysis of dengue virus-specific responses supports an HLA-linked protective role for CD8 ⁺ T cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E2046-53.	3.3	524
40	Memory T Cells in Latent Mycobacterium tuberculosis Infection Are Directed against Three Antigenic Islands and Largely Contained in a CXCR3+CCR6+ Th1 Subset. <i>PLoS Pathogens</i> , 2013, 9, e1003130.	2.1	258
41	Properties of MHC Class I Presented Peptides That Enhance Immunogenicity. <i>PLoS Computational Biology</i> , 2013, 9, e1003266.	1.5	636
42	Immune epitope database analysis resource. <i>Nucleic Acids Research</i> , 2012, 40, W525-W530.	6.5	446
43	Functional classification of class II human leukocyte antigen (HLA) molecules reveals seven different supertypes and a surprising degree of repertoire sharing across supertypes. <i>Immunogenetics</i> , 2011, 63, 325-335.	1.2	351
44	TiArA: A Virtual Appliance for the Analysis of Tiling Array Data. <i>PLoS ONE</i> , 2010, 5, e9993.	1.1	0
45	Pre-existing immunity against swine-origin H1N1 influenza viruses in the general human population. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 20365-20370.	3.3	298
46	Reply to Satheshkumar and Moss: Poxvirus transcriptome analysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, E63-E64.	3.3	4
47	Kinetic Analysis of a Complete Poxvirus Transcriptome Reveals a Novel Class of Genes. <i>FASEB Journal</i> , 2008, 22, .	0.2	0
48	Towards a consensus on datasets and evaluation metrics for developing B-cell epitope prediction tools. <i>Journal of Molecular Recognition</i> , 2007, 20, 75-82.	1.1	209
49	Overcoming the Ontology Enrichment Bottleneck with Quick Term Templates. <i>Nature Precedings</i> , 0, , .	0.1	6