Jack Gorski

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2476255/publications.pdf

Version: 2024-02-01

759233 940533 20 493 12 16 citations h-index g-index papers 20 20 20 408 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Molecular analysis of T cell repertoires. Human Immunology, 1995, 44, 28-34.	2.4	85
2	A Fractal Clonotype Distribution in the CD8+ Memory T Cell Repertoire Could Optimize Potential for Immune Responses. Journal of Immunology, 2003, 170, 3994-4001.	0.8	85
3	A clonotype nomenclature for T cell receptors. Immunogenetics, 2009, 61, 493-502.	2.4	48
4	Complex T Cell Memory Repertoires Participate in Recall Responses at Extremes of Antigenic Load. Journal of Immunology, 2006, 177, 2006-2014.	0.8	35
5	Thymocyte Maturation: Selection for In-Frame TCR α-Chain Rearrangement Is Followed by Selection for Shorter TCR β-Chain Complementarity-Determining Region 3. Journal of Immunology, 2000, 165, 3706-3712.	0.8	33
6	Simulation Studies for a Multistage Dynamic Process of Immune Memory Response to Influenza: Experiment <i>in silico</i> . Annales Zoologici Fennici, 2008, 45, 369-384.	0.6	31
7	Two Compensatory Pathways Maintain Long-Term Stability and Diversity in CD8 T Cell Memory Repertoires. Journal of Immunology, 2009, 183, 2851-2858.	0.8	31
8	Selective T Cell Expansion during Aging of CD8 Memory Repertoires to Influenza Revealed by Modeling. Journal of Immunology, 2011, 186, 6617-6624.	0.8	31
9	Naive T Cell Repertoire Skewing in HLA-A2 Individuals by a Specialized Rearrangement Mechanism Results in Public Memory Clonotypes. Journal of Immunology, 2011, 186, 2970-2977.	0.8	27
10	The Polyclonal CD8 T Cell Response to Influenza M158–66 Generates a Fully Connected Network of Cross-Reactive Clonotypes to Structurally Related Peptides: A Paradigm for Memory Repertoire Coverage of Novel Epitopes or Escape Mutants. Journal of Immunology, 2011, 186, 6390-6397.	0.8	18
11	The functional CD8 T cell memory recall repertoire responding to the influenza A M158–66 epitope is polyclonal and shows a complex clonotype distribution. Human Immunology, 2013, 74, 809-817.	2.4	15
12	CDR3 motif generation and selection in the BV19-utilizing subset of the human CD8 T cell repertoire. Molecular Immunology, 2016, 72, 57-64.	2.2	15
13	CDR3 clonotype and amino acid motif diversity of BV19 expressing circulating human CD8 T cells. Human Immunology, 2016, 77, 137-145.	2.4	14
14	Evidence for preferred MHC class II-TCR recognition independent of the source of bound peptide. European Journal of Immunology, 2002, 32, 2179.	2.9	8
15	Crossâ€reactive responses to modified <scp>M</scp> 1 _{58â€"66} peptides by <scp>CD</scp> 8 ⁺ <scp>T</scp> cells that use noncanonical <scp>BV</scp> genes can describe unknown repertoires. European Journal of Immunology, 2012, 42, 3001-3008.	2.9	8
16	Age-Based Dynamics of a Stable Circulating Cd8 T Cell Repertoire Component. Frontiers in Immunology, 2019, 10, 1717.	4.8	5
17	Measuring Immunological Age: From T Cell Repertoires to Populations. , 2018, , 1-62.		4
18	Structural and Mechanistic Implications of Rearrangement Frequencies within Human TCRBV Genes. Journal of Immunology, 2017, 199, 1142-1152.	0.8	0

#	Article	IF	CITATIONS
19	Measuring Immunological Age: From T cell Repertoires to Populations. , 2018, , 1-60.		O
20	Measuring Immunological Age: From T Cell Repertoires to Populations. , 2019, , 63-124.		0