

Kannan Natarajan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5408520/publications.pdf>

Version: 2024-02-01

29
papers

1,702
citations

361413

20
h-index

477307

29
g-index

32
all docs

32
docs citations

32
times ranked

1843
citing authors

#	ARTICLE	IF	CITATIONS
1	Chaperones and Catalysts: How Antigen Presentation Pathways Cope With Biological Necessity. <i>Frontiers in Immunology</i> , 2022, 13, 859782.	4.8	14
2	Structures of synthetic nanobody-SARS-CoV-2 receptor-binding domain complexes reveal distinct sites of interaction. <i>Journal of Biological Chemistry</i> , 2021, 297, 101202.	3.4	28
3	Structural and dynamic studies of TAPBPR and Tapasin reveal the mechanism of peptide loading of MHC-I molecules. <i>Current Opinion in Immunology</i> , 2020, 64, 71-79.	5.5	19
4	Cutting Edge: Inhibition of the Interaction of NK Inhibitory Receptors with MHC Class I Augments Antiviral and Antitumor Immunity. <i>Journal of Immunology</i> , 2020, 205, 567-572.	0.8	3
5	Alterations in the HLA-B*57:01 Immunopeptidome by Flucloxacillin and Immunogenicity of Drug-Haptenated Peptides. <i>Frontiers in Immunology</i> , 2020, 11, 629399.	4.8	16
6	Structural aspects of chaperone-mediated peptide loading in the MHC-I antigen presentation pathway. <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 2019, 54, 164-173.	5.2	8
7	Cutting antigenic peptides down to size. <i>Journal of Biological Chemistry</i> , 2019, 294, 18545-18546.	3.4	2
8	Structure and Function of Molecular Chaperones that Govern Immune Peptide Loading. <i>Sub-Cellular Biochemistry</i> , 2019, 93, 321-337.	2.4	3
9	MHC Molecules, T cell Receptors, Natural Killer Cell Receptors, and Viral Immune Evasins—Key Elements of Adaptive and Innate Immunity. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1172, 21-62.	1.6	28
10	The Role of Molecular Flexibility in Antigen Presentation and T Cell Receptor-Mediated Signaling. <i>Frontiers in Immunology</i> , 2018, 9, 1657.	4.8	51
11	Peptide exchange on MHC-I by TAPBPR is driven by a negative allosteric release cycle. <i>Nature Chemical Biology</i> , 2018, 14, 811-820.	8.0	74
12	A transgenic mouse model for HLA-B*57:01-linked abacavir drug tolerance and reactivity. <i>Journal of Clinical Investigation</i> , 2018, 128, 2819-2832.	8.2	47
13	An allosteric site in the T-cell receptor C β 2 domain plays a critical signalling role. <i>Nature Communications</i> , 2017, 8, 15260.	12.8	64
14	Crystal structure of a TAPBPR-MHC I complex reveals the mechanism of peptide editing in antigen presentation. <i>Science</i> , 2017, 358, 1064-1068.	12.6	111
15	Interaction of TAPBPR, a tapasin homolog, with MHC-I molecules promotes peptide editing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E1006-15.	7.1	73
16	The cellular environment regulates in situ kinetics of T cell receptor interaction with peptide major histocompatibility complex. <i>European Journal of Immunology</i> , 2015, 45, 2099-2110.	2.9	37
17	The Structure of Mouse Cytomegalovirus m04 Protein Obtained from Sparse NMR Data Reveals a Conserved Fold of the m02-m06 Viral Immune Modulator Family. <i>Structure</i> , 2014, 22, 1263-1273.	3.3	23
18	Mitochondria play a central role in NLRP3 inflammasome activation (349.1). <i>FASEB Journal</i> , 2014, 28, 349.1.	0.5	1

#	ARTICLE	IF	CITATIONS
19	Crystal Structure of the Murine Cytomegalovirus MHC-I Homolog m144. <i>Journal of Molecular Biology</i> , 2006, 358, 157-171.	4.2	36
20	Binding of the Natural Killer Cell Inhibitory Receptor Ly49A to Its Major Histocompatibility Complex Class I Ligand. <i>Journal of Biological Chemistry</i> , 2002, 277, 1433-1442.	3.4	65
21	Structure and Function of Natural Killer Cell Receptors: Multiple Molecular Solutions to Self, Nonself Discrimination. <i>Annual Review of Immunology</i> , 2002, 20, 853-885.	21.8	305
22	MHC class I recognition by Ly49 natural killer cell receptors. <i>Molecular Immunology</i> , 2002, 38, 1023-1027.	2.2	25
23	Structural basis of MHC class I recognition by natural killer cell receptors. <i>Immunological Reviews</i> , 2001, 181, 52-65.	6.0	64
24	A T cell receptor transgenic model of severe, spontaneous organ-specific autoimmunity. <i>European Journal of Immunology</i> , 2001, 31, 2094-2103.	2.9	86
25	Crystal structure of a lectin-like natural killer cell receptor bound to its MHC class I ligand. <i>Nature</i> , 1999, 402, 623-631.	27.8	247
26	Post-thymectomy autoimmune gastritis: fine specificity and pathogenicity of anti-H/K ATPase- reactive T cells. <i>European Journal of Immunology</i> , 1999, 29, 669-677.	2.9	126
27	Interaction of the NK Cell Inhibitory Receptor Ly49A with H-2Dd. <i>Immunity</i> , 1999, 11, 591-601.	14.3	50
28	Post-thymectomy autoimmune gastritis: fine specificity and pathogenicity of anti-H/K ATPase- reactive T cells. <i>European Journal of Immunology</i> , 1999, 29, 669-677.	2.9	5
29	Three-dimensional structure of H-2Dd complexed with an immunodominant peptide from human immunodeficiency virus envelope glycoprotein 120. <i>Journal of Molecular Biology</i> , 1998, 283, 179-191.	4.2	71