

Marvyn T Koning

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

398
citations

933447

10
h-index

794594

19
g-index

23
all docs

23
docs citations

23
times ranked

987
citing authors

#	ARTICLE	IF	CITATIONS
1	Incidence and clinical relevance of cage subsidence in anterior cervical discectomy and fusion: a systematic review. <i>Acta Neurochirurgica</i> , 2018, 160, 873-880.	1.7	69
2	MuSK myasthenia gravis monoclonal antibodies. <i>Neurology: Neuroimmunology and NeuroInflammation</i> , 2019, 6, e547.	6.0	64
3	<i>I</i> GLV3-21 <i>I</i> *01 is an inherited risk factor for CLL through the acquisition of a single-point mutation enabling autonomous BCR signaling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 4320-4327.	7.1	55
4	B-cell receptor sequencing of anti-citrullinated protein antibody (ACPA) IgG-expressing B cells indicates a selective advantage for the introduction of N-glycosylation sites during somatic hypermutation. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, annrheumdis-2017-212052.	0.9	51
5	ARTISAN PCR: rapid identification of full-length immunoglobulin rearrangements without primer binding bias. <i>British Journal of Haematology</i> , 2017, 178, 983-986.	2.5	28
6	A Comparison of Immunoglobulin Variable Region N-Linked Glycosylation in Healthy Donors, Autoimmune Disease and Lymphoma. <i>Frontiers in Immunology</i> , 2020, 11, 241.	4.8	28
7	N-Glycosylation Site Analysis of Citrullinated Antigen-Specific B-Cell Receptors Indicates Alternative Selection Pathways During Autoreactive B-Cell Development. <i>Frontiers in Immunology</i> , 2019, 10, 2092.	4.8	23
8	Response to: "Acquiring new N-glycosylation sites in variable regions of immunoglobulin genes by somatic hypermutation is a common feature of autoimmune diseases" by Visser et al. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, e70-e70.	0.9	15
9	Template-switching anchored polymerase chain reaction reliably amplifies functional lambda light chain transcripts of malignant lymphoma. <i>Leukemia and Lymphoma</i> , 2014, 55, 1212-1214.	1.3	14
10	Acquired N-Linked Glycosylation Motifs in B-Cell Receptors of Primary Cutaneous B-Cell Lymphoma and the Normal B-Cell Repertoire. <i>Journal of Investigative Dermatology</i> , 2019, 139, 2195-2203.	0.7	12
11	Peripheral IgE Repertoires of Healthy Donors Carry Moderate Mutation Loads and Do Not Overlap With Other Isotypes. <i>Frontiers in Immunology</i> , 2019, 10, 1543.	4.8	10
12	A case of fulminant Epstein-Barr virus encephalitis in an immune-competent adult. <i>Journal of NeuroVirology</i> , 2019, 25, 422-425.	2.1	9
13	Tandem Substitutions in Somatic Hypermutation. <i>Blood</i> , 2021, 138, 996-996.	1.4	6
14	Templated insertions at VD and DJ junctions create unique B-cell receptors in the healthy B-cell repertoire. <i>European Journal of Immunology</i> , 2020, 50, 2099-2101.	2.9	3
15	Primary Cutaneous Follicle Center Lymphomas (PCFCL) Express Heavily Mutated B-Cell Receptors with Acquired N-Glycosylation Motifs and Lack Ongoing Somatic Hypermutation. <i>Blood</i> , 2018, 132, 1573-1573.	1.4	3
16	Primary Mediastinal Large B-Cell Lymphoma Exhibits Autonomous BCR Signaling and Responds to the Second Generation BTK Inhibitor Acalabrutinib. <i>Blood</i> , 2016, 128, 4171-4171.	1.4	2
17	Tandem Substitutions in Somatic Hypermutation. <i>Frontiers in Immunology</i> , 2021, 12, 807015.	4.8	2
18	Two Independent Hematological malignancies in a B-Cell Deficient Good Syndrome Patient. <i>Rheumatology</i> , 2021, 60, e126-e128.	1.9	1

#	ARTICLE	IF	CITATIONS
19	Endogenous Immunoglobulin-Derived Neopeptides Are Processed and Form a Sizeable Fraction of the HLA Class I Ligandome of Human Lymphoma Cells. <i>Blood</i> , 2016, 128, 914-914.	1.4	1
20	Antigen-Independent, Autonomous B-Cell Receptor Signaling As a Dominant Candidate Oncogenic Mechanism in ABC DLBCL. <i>Blood</i> , 2016, 128, 778-778.	1.4	1
21	High-Throughput BCR Sequencing and Single-Cell Transcriptomics Reveal Distinct Transcriptional Profiles Associated with Subclonal Evolution of Follicular Lymphoma. <i>Blood</i> , 2019, 134, 298-298.	1.4	1
22	Massive Parallel Sequencing of Full-Length B-Cell Receptor Sequences Reveals HLA-Dependent Shaping of the B-Cell Immune Repertoire. <i>Blood</i> , 2014, 124, 4143-4143.	1.4	0
23	The B-Cell Receptor of Primary Cutaneous Follicle Center Lymphoma: Implications for Pathogenesis. <i>Blood</i> , 2016, 128, 4136-4136.	1.4	0