

Bertrand Nadel

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

2,333
citations

293460

24
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340414

39
g-index

42
all docs

42
docs citations

42
times ranked

3248
citing authors

#	ARTICLE	IF	CITATIONS
1	The EHA Research Roadmap: Malignant Lymphoid Diseases. <i>HemaSphere</i> , 2022, 6, e726.	1.2	1
2	Follicular lymphoma dynamics. <i>Advances in Immunology</i> , 2021, 150, 43-103.	1.1	19
3	The Premalignant Ancestor Cell of t(14;18)+ Lymphoma. <i>HemaSphere</i> , 2021, 5, e579.	1.2	5
4	MYC deficiency impairs the development of effector/memory T lymphocytes. <i>IScience</i> , 2021, 24, 102761.	1.9	10
5	Human B Lymphomas Reveal Their Secrets Through Genetic Mouse Models. <i>Frontiers in Immunology</i> , 2021, 12, 683597.	2.2	6
6	Recurrent Crebbp Mutations in Follicular Lymphoma Appear Localized to the Committed B-Cell Lineage. <i>Blood</i> , 2020, 136, 30-31.	0.6	2
7	Follicular lymphoma: Biology update. <i>Hematologie</i> , 2019, 25, 23-31.	0.0	0
8	Single-Cell RNA Sequencing Identifies a Pseudo-Immune Differentiation Axis As the Main Source of Functional Heterogeneity in Follicular Lymphoma B-Cells. <i>Blood</i> , 2019, 134, 548-548.	0.6	6
9	Fit $\hat{1}^2$ T-cell receptor suppresses leukemogenesis of Pten-deficient thymocytes. <i>Haematologica</i> , 2018, 103, 999-1007.	1.7	6
10	Human germinal center transcriptional programs are de-synchronized in B cell lymphoma. <i>Nature Immunology</i> , 2018, 19, 1013-1024.	7.0	115
11	Desynchronization of the Germinal Center Dynamics and Remodeling of the Tumor Microenvironment Characterize KMT2D-Driven Lymphomagenesis. <i>Blood</i> , 2018, 132, 670-670.	0.6	8
12	Follicular lymphoma: State-of-the-art ICML workshop in Lugano 2015. <i>Hematological Oncology</i> , 2017, 35, 397-407.	0.8	11
13	BCL-B (BCL2L10) is overexpressed in patients suffering from multiple myeloma (MM) and drives an MM-like disease in transgenic mice. <i>Journal of Experimental Medicine</i> , 2016, 213, 1705-1722.	4.2	24
14	Contiguous follicular lymphoma and follicular lymphoma in situ harboring N-glycosylated sites. <i>Haematologica</i> , 2015, 100, e155-e157.	1.7	17
15	Premalignant cell dynamics in indolent B-cell malignancies. <i>Current Opinion in Hematology</i> , 2015, 22, 388-396.	1.2	13
16	Nature and importance of follicular lymphoma precursors. <i>Haematologica</i> , 2014, 99, 802-810.	1.7	47
17	Human t(14;18)positive germinal center B cells: a new step in follicular lymphoma pathogenesis?. <i>Blood</i> , 2014, 123, 3462-3465.	0.6	44
18	Early lesions of follicular lymphoma: a genetic perspective. <i>Haematologica</i> , 2014, 99, 481-488.	1.7	91

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19	t(14;18) Translocation: A Predictive Blood Biomarker for Follicular Lymphoma. <i>Journal of Clinical Oncology</i> , 2014, 32, 1347-1355.	0.8	115
20	Germinal center reentries of BCL2-overexpressing B cells drive follicular lymphoma progression. <i>Journal of Clinical Investigation</i> , 2014, 124, 5337-5351.	3.9	96
21	Therapeutic Targeting of c-Myc in T-Cell Acute Lymphoblastic Leukemia (T-ALL). <i>Oncotarget</i> , 2014, 5, 3168-3172.	0.8	58
22	Toward a <i>NOTCH1/FBXW7/RAS/PTEN</i> -Based Oncogenetic Risk Classification of Adult T-Cell Acute Lymphoblastic Leukemia: A Group for Research in Adult Acute Lymphoblastic Leukemia Study. <i>Journal of Clinical Oncology</i> , 2013, 31, 4333-4342.	0.8	202
23	Extensive molecular mapping of <i>TCRα</i> - and <i>TCRβ</i> -involved chromosomal translocations reveals distinct mechanisms of oncogene activation in T-ALL. <i>Blood</i> , 2012, 120, 3298-3309.	0.6	31
24	TLX Homeodomain Oncogenes Mediate T Cell Maturation Arrest in T-ALL via Interaction with ETS1 and Suppression of <i>TCRα</i> Gene Expression. <i>Cancer Cell</i> , 2012, 21, 563-576.	7.7	81
25	Early Steps of Follicular Lymphoma Pathogenesis. <i>Advances in Immunology</i> , 2011, 111, 1-46.	1.1	91
26	Posttranscriptional deregulation of MYC via PTEN constitutes a major alternative pathway of MYC activation in T-cell acute lymphoblastic leukemia. <i>Blood</i> , 2011, 117, 6650-6659.	0.6	72
27	Agricultural pesticide exposure and the molecular connection to lymphomagenesis. <i>Journal of Experimental Medicine</i> , 2009, 206, 1473-1483.	4.2	73
28	V(D)J targeting mistakes occur at low frequency in acute lymphoblastic leukemia. <i>Genes Chromosomes and Cancer</i> , 2009, 48, 725-736.	1.5	9
29	Pathophysiological aspects of memory B-cell development. <i>Trends in Immunology</i> , 2008, 29, 25-33.	2.9	33
30	The C-MYB locus is involved in chromosomal translocation and genomic duplications in human T-cell acute leukemia (T-ALL), the translocation defining a new T-ALL subtype in very young children. <i>Blood</i> , 2007, 110, 1251-1261.	0.6	249
31	Different chromosomal breakpoints impact the level of LMO2 expression in T-ALL. <i>Blood</i> , 2007, 110, 388-392.	0.6	47
32	In Vivo Reinsertion of Excised Episomes by the V(D)J Recombinase: A Potential Threat to Genomic Stability. <i>PLoS Biology</i> , 2007, 5, e43.	2.6	31
33	Recombinase, chromosomal translocations and lymphoid neoplasia: Targeting mistakes and repair failures. <i>DNA Repair</i> , 2006, 5, 1246-1258.	1.3	90
34	Follicular lymphoma-like B cells in healthy individuals: a novel intermediate step in early lymphomagenesis. <i>Journal of Experimental Medicine</i> , 2006, 203, 2425-2431.	4.2	187
35	Unraveling the Consecutive Recombination Events in the Human <i>IGK</i> Locus. <i>Journal of Immunology</i> , 2004, 173, 3878-3888.	0.4	28
36	Distinct t(7;9)(q34;q32) breakpoints in healthy individuals and individuals with T-ALL. <i>Nature Genetics</i> , 2003, 33, 342-344.	9.4	26

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37	V(D)J-mediated Translocations in Lymphoid Neoplasms: A Functional Assessment of Genomic Instability by Cryptic Sites. <i>Journal of Experimental Medicine</i> , 2002, 195, 85-98.	4.2	150
38	Novel Insights into the Mechanism of t(14;18)(q32;q21) Translocation in Follicular Lymphoma. <i>Leukemia and Lymphoma</i> , 2001, 42, 1181-1194.	0.6	17
39	Follicular lymphomas' BCL-2/IgH junctions contain templated nucleotide insertions: novel insights into the mechanism of t(14;18) translocation. <i>Blood</i> , 2000, 95, 3520-3529.	0.6	142
40	Sequence of the Spacer in the Recombination Signal Sequence Affects V(D)J Rearrangement Frequency and Correlates with Nonrandom V _H Usage In Vivo. <i>Journal of Experimental Medicine</i> , 1998, 187, 1495-1503.	4.2	77