Sophia Yancopoulos

List of Publications by Year in descending order

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933447 1372567 12 995 10 10 citations g-index h-index papers 13 13 13 1346 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Myeloid-derived suppressor cell subtypes differentially influence T-cell function, T-helper subset differentiation, and clinical course in CLL. Leukemia, 2021, 35, 3163-3175.	7.2	25
2	A seven-gene expression panel distinguishing clonal expansions of pre-leukemic and chronic lymphocytic leukemia B cells from normal B lymphocytes. Immunologic Research, 2015, 63, 90-100.	2.9	18
3	Autoantigen can promote progression to a more aggressive TCL1 leukemia by selecting variants with enhanced B-cell receptor signaling. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E1500-7.	7.1	49
4	Intraclonal Complexity in Chronic Lymphocytic Leukemia: Fractions Enriched in Recently Born/Divided and Older/Quiescent Cells. Molecular Medicine, 2011, 17, 1374-1382.	4.4	140
5	Identification of outcome-correlated cytokine clusters in chronic lymphocytic leukemia. Blood, 2011, 118, 5201-5210.	1.4	110
6	The rise and fall of breakpoint reuse depending on genome resolution. BMC Bioinformatics, 2011, 12, S1.	2.6	19
7	DCJ Path Formulation for Genome Transformations which Include Insertions, Deletions, and Duplications. Journal of Computational Biology, 2009, 16, 1311-1338.	1.6	32
8	Genome Rearrangement by the Double Cut and Join Operation. Methods in Molecular Biology, 2008, 452, 385-416.	0.9	13
9	B cell receptors in TCL1 transgenic mice resemble those of aggressive, treatment-resistant human chronic lymphocytic leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 11713-11718.	7.1	154
10	IGHV Gene Replacement in B-Cell Chronic Lymphocytic Leukemia (B-CLL) Occurs at a Frequency Similar to That in Normal B Cells and May Augment Clonal Expansion by Permitting Autogenic/Microbial Clonal Stimulation Blood, 2006, 108, 2086-2086.	1.4	0
11	Genome Analysis of CLL by Representational Oligonucleotide Microarray Analysis (ROMA) Blood, 2006, 108, 2085-2085.	1.4	O
12	Efficient sorting of genomic permutations by translocation, inversion and block interchange. Bioinformatics, 2005, 21, 3340-3346.	4.1	435