Eugene M Oltz

List of Publications by Year in descending order

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FUCENE MOITZ

#	Article	IF	CITATIONS
1	Gene Regulatory Circuits in Innate and Adaptive Immune Cells. Annual Review of Immunology, 2022, 40, 387-411.	21.8	6
2	SARS-CoV-2 spreads through cell-to-cell transmission. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	145
3	Whole-genome profiling of DNA methylation and hydroxymethylation identifies distinct regulatory programs among innate lymphocytes. Nature Immunology, 2022, 23, 619-631.	14.5	14
4	Neutralizing antibody responses elicited by SARS-CoV-2 mRNA vaccination wane over time and are boosted by breakthrough infection. Science Translational Medicine, 2022, 14, eabn8057.	12.4	150
5	Neutralization of SARS-CoV-2 Omicron sub-lineages BA.1, BA.1.1, and BA.2. Cell Host and Microbe, 2022, 30, 1093-1102.e3.	11.0	114
6	Caspase-4/11 exacerbates disease severity in SARS–CoV-2 infection by promoting inflammation and immunothrombosis. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2202012119.	7.1	25
7	Enhanced epigenetic profiling of classical human monocytes reveals a specific signature of healthy aging in the DNA methylome. Nature Aging, 2021, 1, 124-141.	11.6	30
8	Cancer-associated exportin-6 upregulation inhibits the transcriptionally repressive and anticancer effects of nuclear profilin-1. Cell Reports, 2021, 34, 108749.	6.4	9
9	Loss of synergistic transcriptional feedback loops drives diverse B-cell cancers. EBioMedicine, 2021, 71, 103559.	6.1	1
10	Neutralization of SARS-CoV-2 Variants of Concern Harboring Q677H. MBio, 2021, 12, e0251021.	4.1	33
11	Impaired neutralizing antibody response to COVID-19 mRNA vaccines in cancer patients. Cell and Bioscience, 2021, 11, 197.	4.8	32
12	Barrier-to-Autointegration Factor 1 Protects against a Basal cGAS-STING Response. MBio, 2020, 11, .	4.1	33
13	DNA double-strand breaks induce H2Ax phosphorylation domains in a contact-dependent manner. Nature Communications, 2020, 11, 3158.	12.8	97
14	Blood natural killer cell deficiency reveals an immunotherapy strategy for atopic dermatitis. Science Translational Medicine, 2020, 12, .	12.4	57
15	Neutralizing antibody against SARS-CoV-2 spike in COVID-19 patients, health care workers, and convalescent plasma donors. JCI Insight, 2020, 5, .	5.0	86
16	Circadian rhythm–dependent and circadian rhythm–independent impacts of the molecular clock on type 3 innate lymphoid cells. Science Immunology, 2019, 4, .	11.9	65
17	Subsets of ILC3â^'ILC1-like cells generate a diversity spectrum of innate lymphoid cells in human mucosal tissues. Nature Immunology, 2019, 20, 980-991.	14.5	141
18	Regional Gene Repression by DNA Double-Strand Breaks in G ₁ Phase Cells. Molecular and Cellular Biology, 2019, 39, .	2.3	15

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19	Gene Regulatory Programs Conferring Phenotypic Identities to Human NK Cells. Cell, 2019, 176, 348-360.e12.	28.9	125
20	A B-Cell-Specific Enhancer Orchestrates Nuclear Architecture to Generate a Diverse Antigen Receptor Repertoire. Molecular Cell, 2019, 73, 48-60.e5.	9.7	29
21	Toxoplasma gondii infection drives conversion of NK cells into ILC1-like cells. ELife, 2019, 8, .	6.0	91
22	Targeted epigenetic repression of a lymphoma oncogene by sequence-specific histone modifiers induces apoptosis in DLBCL. Leukemia and Lymphoma, 2017, 58, 445-456.	1.3	5
23	cis -Regulatory Circuits Regulating NEK6 Kinase Overexpression in Transformed B Cells Are Super-Enhancer Independent. Cell Reports, 2017, 18, 2918-2931.	6.4	14
24	Activation of Mouse <i>Tcrb</i> : Uncoupling RUNX1 Function from Its Cooperative Binding with ETS1. Journal of Immunology, 2017, 199, 1131-1141.	0.8	10
25	NKG2D–NKG2D Ligand Interaction Inhibits the Outgrowth of Naturally Arising Low-Grade B Cell Lymphoma In Vivo. Journal of Immunology, 2016, 196, 4805-4813.	0.8	19
26	Distinct Gene Regulatory Pathways for Human Innate versus Adaptive Lymphoid Cells. Cell, 2016, 165, 1134-1146.	28.9	134
27	The Colonic Crypt Protects Stem Cells from Microbiota-Derived Metabolites. Cell, 2016, 165, 1708-1720.	28.9	484
28	Mapping of Variable DNA Methylation Across Multiple Cell Types Defines a Dynamic Regulatory Landscape of the Human Genome. G3: Genes, Genomes, Genetics, 2016, 6, 973-986.	1.8	41
29	CYLD and the NEMO Zinc Finger Regulate Tumor Necrosis Factor Signaling and Early Embryogenesis. Journal of Biological Chemistry, 2015, 290, 22076-22084.	3.4	11
30	Domain-Specific and Stage-Intrinsic Changes in <i>Tcrb</i> Conformation during Thymocyte Development. Journal of Immunology, 2015, 195, 1262-1272.	0.8	11
31	Regulation of Tcrb Gene Assembly by Genetic, Epigenetic, and Topological Mechanisms. Advances in Immunology, 2015, 128, 273-306.	2.2	27
32	Enhancer Sequence Variants and Transcription-Factor Deregulation Synergize to Construct Pathogenic Regulatory Circuits in B-Cell Lymphoma. Immunity, 2015, 42, 186-198.	14.3	64
33	Lineage-specific compaction of <i>Tcrb</i> requires a chromatin barrier to protect the function of a long-range tethering element. Journal of Experimental Medicine, 2015, 212, 107-120.	8.5	54
34	The histone methyltransferase SETDB1 represses endogenous and exogenous retroviruses in B lymphocytes. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 8367-8372.	7.1	78
35	Short-Circuiting Gene Regulatory Networks: Origins of B Cell Lymphoma. Trends in Genetics, 2015, 31, 720-731.	6.7	5
36	HCoDES Reveals Chromosomal DNA End Structures with Single-Nucleotide Resolution. Molecular Cell, 2014, 56, 808-818.	9.7	31

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37	Functional Intersection of ATM and DNA-Dependent Protein Kinase Catalytic Subunit in Coding End Joining during V(D)J Recombination. Molecular and Cellular Biology, 2013, 33, 3568-3579.	2.3	39
38	Unifying model for molecular determinants of the preselection Vβ repertoire. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E3206-15.	7.1	50
39	Defining the Malignant Epigenome in Non-Hodgkin Lymphoma. Blood, 2012, 120, 524-524.	1.4	1