

Peter van Galen

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

5,034
citations

411340

20
h-index

563245

28
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37
docs citations

37
times ranked

11397
citing authors

#	ARTICLE	IF	CITATIONS
1	Association of clonal hematopoiesis with chronic obstructive pulmonary disease. <i>Blood</i> , 2022, 139, 357-368.	0.6	106
2	Renewing your HBO1 subscription. <i>Blood</i> , 2022, 139, 802-804.	0.6	1
3	Mitochondrial variant enrichment from high-throughput single-cell RNA sequencing resolves clonal populations. <i>Nature Biotechnology</i> , 2022, 40, 1030-1034.	9.4	45
4	Single-Cell Multiomics Reveals Clonal T-Cell Expansions and Exhaustion in Blastic Plasmacytoid Dendritic Cell Neoplasm. <i>Frontiers in Immunology</i> , 2022, 13, 809414.	2.2	6
5	Plasticity and immune evasion in childhood ALL. <i>Blood</i> , 2022, 139, 2096-2097.	0.6	0
6	A cellular hierarchy framework for understanding heterogeneity and predicting drug response in acute myeloid leukemia. <i>Nature Medicine</i> , 2022, 28, 1212-1223.	15.2	104
7	WAT3R: recovery of T-cell receptor variable regions from 3â€² single-cell RNA-sequencing. <i>Bioinformatics</i> , 2022, 38, 3645-3647.	1.8	2
8	Clonal barcoding with qPCR detection enables live cell functional analyses for cancer research. <i>Nature Communications</i> , 2022, 13, .	5.8	1
9	Single-Cell RNA Sequencing to Disentangle the Blood System. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2021, 41, 1012-1018.	1.1	8
10	Extended-representation bisulfite sequencing of gene regulatory elements in multiplexed samples and single cells. <i>Nature Biotechnology</i> , 2021, 39, 1086-1094.	9.4	28
11	Single-cell lineage analysis reveals genetic and epigenetic interplay in glioblastoma drug resistance. <i>Genome Biology</i> , 2020, 21, 174.	3.8	59
12	Decoding the Noncoding Cancer Genome. <i>Cancer Discovery</i> , 2020, 10, 646-647.	7.7	2
13	Chromatin accessibility promotes hematopoietic and leukemia stem cell activity. <i>Nature Communications</i> , 2020, 11, 1406.	5.8	32
14	A novel method for detecting the cellular stemness state in normal and leukemic human hematopoietic cells can predict disease outcome and drug sensitivity. <i>Leukemia</i> , 2019, 33, 2061-2077.	3.3	13
15	A stemness screen reveals C3orf54/INKA1 as a promoter of human leukemia stem cell latency. <i>Blood</i> , 2019, 133, 2198-2211.	0.6	25
16	Single-Cell RNA-Seq Reveals AML Hierarchies Relevant to Disease Progression and Immunity. <i>Cell</i> , 2019, 176, 1265-1281.e24.	13.5	642
17	Developmental and oncogenic programs in H3K27M gliomas dissected by single-cell RNA-seq. <i>Science</i> , 2018, 360, 331-335.	6.0	461
18	Dual Targeting of Oncogenic Activation and Inflammatory Signaling Increases Therapeutic Efficacy in Myeloproliferative Neoplasms. <i>Cancer Cell</i> , 2018, 33, 29-43.e7.	7.7	186

#	ARTICLE	IF	CITATIONS
19	Integrated Stress Response Activity Marks Stem Cells in Normal Hematopoiesis and Leukemia. <i>Cell Reports</i> , 2018, 25, 1109-1117.e5.	2.9	88
20	Single-Cell RNA-Seq Reveals AML Cellular Hierarchies Relevant to Clinical Outcomes and Immunity. <i>Blood</i> , 2018, 132, 542-542.	0.6	0
21	Identification of Gene Regulatory Networks Governing Stemness Properties of Human HSC and LSC. <i>Blood</i> , 2018, 132, 3832-3832.	0.6	0
22	Adaptive Chromatin Remodeling Drives Glioblastoma Stem Cell Plasticity and Drug Tolerance. <i>Cell Stem Cell</i> , 2017, 20, 233-246.e7.	5.2	387
23	A Multiplexed System for Quantitative Comparisons of Chromatin Landscapes. <i>Molecular Cell</i> , 2016, 61, 170-180.	4.5	111
24	EHMT1 and EHMT2 inhibition induces fetal hemoglobin expression. <i>Blood</i> , 2015, 126, 1930-1939.	0.6	76
25	Two new routes to make blood: Hematopoietic specification from pluripotent cell lines versus reprogramming of somatic cells. <i>Experimental Hematology</i> , 2015, 43, 756-759.	0.2	5
26	The unfolded protein response governs integrity of the haematopoietic stem-cell pool during stress. <i>Nature</i> , 2014, 510, 268-272.	13.7	292
27	Self-renewal as a therapeutic target in human colorectal cancer. <i>Nature Medicine</i> , 2014, 20, 29-36.	15.2	438
28	Reduced Lymphoid Lineage Priming Promotes Human Hematopoietic Stem Cell Expansion. <i>Cell Stem Cell</i> , 2014, 14, 94-106.	5.2	63
29	Variable Clonal Repopulation Dynamics Influence Chemotherapy Response in Colorectal Cancer. <i>Science</i> , 2013, 339, 543-548.	6.0	691
30	Attenuation of miR-126 Activity Expands HSC In Vivo without Exhaustion. <i>Cell Stem Cell</i> , 2012, 11, 799-811.	5.2	197
31	Stem cell gene expression programs influence clinical outcome in human leukemia. <i>Nature Medicine</i> , 2011, 17, 1086-1093.	15.2	894
32	Rise of the Clones: Myelodysplastic Syndrome  to Secondary Acute Myeloid Leukemia. <i>Blood Cancer Discovery</i> , 0, , OF1-OF3.	2.6	0