

Jake A Kloeber

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5576526/publications.pdf>

Version: 2024-02-01

29
papers

594
citations

687363

13
h-index

713466

21
g-index

30
all docs

30
docs citations

30
times ranked

1028
citing authors

#	ARTICLE	IF	CITATIONS
1	Critical DNA damaging pathways in tumorigenesis. <i>Seminars in Cancer Biology</i> , 2022, 85, 164-184.	9.6	8
2	Cell-free DNA for the detection of emerging treatment failure in relapsed/ refractory multiple myeloma. <i>Leukemia</i> , 2022, 36, 1078-1087.	7.2	13
3	m6A demethylation of cytidine deaminase APOBEC3B mRNA orchestrates arsenic-induced mutagenesis. <i>Journal of Biological Chemistry</i> , 2022, 298, 101563.	3.4	10
4	The deubiquitinase USP7 regulates oxidative stress through stabilization of HO-1. <i>Oncogene</i> , 2022, 41, 4018-4027.	5.9	8
5	Single-cell RNA-seq reveals developmental plasticity with coexisting oncogenic states and immune evasion programs in ETP-ALL. <i>Blood</i> , 2021, 137, 2463-2480.	1.4	35
6	LRRK2 inhibition potentiates PARP inhibitor cytotoxicity through inhibiting homologous recombination-mediated DNA double strand break repair. <i>Clinical and Translational Medicine</i> , 2021, 11, e341.	4.0	7
7	USP13 regulates the replication stress response by deubiquitinating TopBP1. <i>DNA Repair</i> , 2021, 100, 103063.	2.8	10
8	MET Amplification Attenuates Lung Tumor Response to Immunotherapy by Inhibiting STING. <i>Cancer Discovery</i> , 2021, 11, 2726-2737.	9.4	35
9	ASTE1 promotes shieldin-complex-mediated DNA repair by attenuating end resection. <i>Nature Cell Biology</i> , 2021, 23, 894-904.	10.3	28
10	Single-Cell Profiling Reveals Metabolic Reprogramming as a Resistance Mechanism in <i>BRAF</i> -Mutated Multiple Myeloma. <i>Clinical Cancer Research</i> , 2021, 27, 6432-6444.	7.0	18
11	Dynamic transcriptional reprogramming leads to immunotherapeutic vulnerabilities in myeloma. <i>Nature Cell Biology</i> , 2021, 23, 1199-1211.	10.3	22
12	RNF19A-mediated ubiquitination of BARD1 prevents BRCA1/BARD1-dependent homologous recombination. <i>Nature Communications</i> , 2021, 12, 6653.	12.8	7
13	Maturity State and MCL-1 Dependence Predetermines Response to NOTCH1 Inhibition in T-ALL. <i>Blood</i> , 2021, 138, 3484-3484.	1.4	0
14	FoxM1 insufficiency hyperactivates Ect2-RhoA-Dia1 signaling to drive cancer. <i>Nature Cancer</i> , 2020, 1, 1010-1024.	13.2	6
15	The deubiquitinase USP36 Regulates DNA replication stress and confers therapeutic resistance through PrimPol stabilization. <i>Nucleic Acids Research</i> , 2020, 48, 12711-12726.	14.5	26
16	USP52 regulates DNA end resection and chemosensitivity through removing inhibitory ubiquitination from CtIP. <i>Nature Communications</i> , 2020, 11, 5362.	12.8	16
17	DNA end resection and its role in DNA replication and DSB repair choice in mammalian cells. <i>Experimental and Molecular Medicine</i> , 2020, 52, 1705-1714.	7.7	72
18	Eosinophil-derived IL-13 promotes emphysema. <i>European Respiratory Journal</i> , 2019, 53, 1801291.	6.7	47

#	ARTICLE	IF	CITATIONS
19	Determining resistance mechanisms in BRAF-mutated multiple myeloma. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, e22.	0.4	0
20	Circulating Tumor DNA in the Peripheral Blood As Early Predictor of Clinical Outcome in Relapsed/Refractory Multiple Myeloma. <i>Blood</i> , 2019, 134, 4350-4350.	1.4	0
21	Determining Resistance Mechanisms in BRAF-mutated Multiple Myeloma. <i>Blood</i> , 2019, 134, 316-316.	1.4	0
22	Single Cell RNA-Seq Reveals Deranged Developmental Hierarchy with Coexisting Oncogenic States and Immune Evasion Programs in ETP T-ALL. <i>Blood</i> , 2019, 134, 3953-3953.	1.4	0
23	Enhancer Rewiring Dependent Switch from BCL2 to MCL1 Dependency Predicts NOTCH1 Inhibition Response in T-ALL. <i>Blood</i> , 2019, 134, 3948-3948.	1.4	0
24	Defining the Differentiation States of Multiple Myeloma at Single Cell Resolution Reveals Opportunities for Immunotherapy. <i>Blood</i> , 2019, 134, 3091-3091.	1.4	0
25	Immunological Consequences of Lenalidomide with and without Dexamethasone in Newly Diagnosed Multiple Myeloma. <i>Blood</i> , 2019, 134, 3070-3070.	1.4	0
26	Genomic discovery and clonal tracking in multiple myeloma by cell-free DNA sequencing. <i>Leukemia</i> , 2018, 32, 1838-1841.	7.2	42
27	Eosinophil-dependent skin innervation and itching following contact toxicant exposure in mice. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 135, 477-487.e1.	2.9	31
28	Homologous recombination into the eosinophil peroxidase locus generates a strain of mice expressing Cre recombinase exclusively in eosinophils. <i>Journal of Leukocyte Biology</i> , 2013, 94, 17-24.	3.3	85
29	Expression of the secondary granule proteins major basic protein 1 (MBP-1) and eosinophil peroxidase (EPX) is required for eosinophilopoiesis in mice. <i>Blood</i> , 2013, 122, 781-790.	1.4	68