Tamir Chandra

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6192846/publications.pdf

Version: 2024-02-01

361296 477173 4,377 29 20 29 citations h-index g-index papers 37 37 37 8266 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	SC3: consensus clustering of single-cell RNA-seq data. Nature Methods, 2017, 14, 483-486.	9.0	1,203
2	cGAS surveillance of micronuclei links genome instability to innate immunity. Nature, 2017, 548, 461-465.	13.7	1,158
3	Global Reorganization of the Nuclear Landscape in Senescent Cells. Cell Reports, 2015, 10, 471-483.	2.9	282
4	Redistribution of the Lamin B1 genomic binding profile affects rearrangement of heterochromatic domains and SAHF formation during senescence. Genes and Development, 2013, 27, 1800-1808.	2.7	259
5	Independence of Repressive Histone Marks and Chromatin Compaction during Senescent Heterochromatic Layer Formation. Molecular Cell, 2012, 47, 203-214.	4.5	258
6	Partial reprogramming induces a steady decline in epigenetic age before loss of somatic identity. Aging Cell, 2019, 18, e12877.	3.0	128
7	An epigenome-wide association study of sex-specific chronological ageing. Genome Medicine, 2020, 12, 1.	3.6	117
8	Experimental design for single-cell RNA sequencing. Briefings in Functional Genomics, 2018, 17, 233-239.	1.3	113
9	Kidney Single-Cell Atlas Reveals Myeloid Heterogeneity in Progression and Regression of Kidney Disease. Journal of the American Society of Nephrology: JASN, 2020, 31, 2833-2854.	3.0	113
10	Notch Signaling Mediates Secondary Senescence. Cell Reports, 2019, 27, 997-1007.e5.	2.9	82
11	Epigenetic age prediction. Aging Cell, 2021, 20, e13452.	3.0	81
12	Proliferation Drives Aging-Related Functional Decline in a Subpopulation of the Hematopoietic Stem Cell Compartment. Cell Reports, 2017, 19, 1503-1511.	2.9	76
13	Polymer Modeling Predicts Chromosome Reorganization in Senescence. Cell Reports, 2019, 28, 3212-3223.e6.	2.9	60
14	Cellular reprogramming and epigenetic rejuvenation. Clinical Epigenetics, 2021, 13, 170.	1.8	54
15	Multi-layered Spatial Transcriptomics Identify Secretory Factors Promoting Human Hematopoietic Stem Cell Development. Cell Stem Cell, 2020, 27, 822-839.e8.	5.2	51
16	Functional heterogeneity in senescence. Biochemical Society Transactions, 2020, 48, 765-773.	1.6	50
17	Phenotype Specific Analyses Reveal Distinct Regulatory Mechanism for Chronically Activated p53. PLoS Genetics, 2015, 11, e1005053.	1.5	47
18	Chromosome organisation during ageing and senescence. Current Opinion in Cell Biology, 2016, 40, 161-167.	2.6	44

#	Article	IF	CITATIONS
19	Age-related clonal haemopoiesis is associated with increased epigenetic age. Current Biology, 2019, 29, R786-R787.	1.8	37
20	Longitudinal dynamics of clonal hematopoiesis identifies gene-specific fitness effects. Nature Medicine, 2022, 28, 1439-1446.	15.2	36
21	Wilms Tumor 1b defines a wound-specific sheath cell subpopulation associated with notochord repair. ELife, 2018, 7, .	2.8	21
22	Clonal haematopoiesis of indeterminate potential: intersections between inflammation, vascular disease and heart failure. Clinical Science, 2021, 135, 991-1007.	1.8	18
23	Inhibition of the 60S ribosome biogenesis GTPase LSG1 causes endoplasmic reticular disruption and cellular senescence. Aging Cell, 2019, 18, e12981.	3.0	17
24	Tfap2b specifies an embryonic melanocyte stem cell that retains adult multifate potential. Cell Reports, 2022, 38, 110234.	2.9	15
25	Purifying stem cellâ€derived red blood cells: a highâ€throughput labelâ€free downstream processing strategy based on microfluidic spiral inertial separation and membrane filtration. Biotechnology and Bioengineering, 2020, 117, 2032-2045.	1.7	13
26	Induction and transmission of oncogene-induced senescence. Cellular and Molecular Life Sciences, 2021, 78, 843-852.	2.4	11
27	A Stem Cell Reporter for Investigating Pluripotency and Self-Renewal in the Rat. Stem Cell Reports, 2020, 14, 154-166.	2.3	6
28	Multiplexing for Oxidative Bisulfite Sequencing (oxBS-seq). Methods in Molecular Biology, 2018, 1708, 665-678.	0.4	5
29	Clonality in haematopoietic stem cell ageing. Mechanisms of Ageing and Development, 2020, 189, 111279.	2.2	4