Jan Schroeder

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/129243/publications.pdf

Version: 2024-02-01

24 papers 1,915 citations

14 h-index

623734

610901 24 g-index

24 all docs

24 docs citations

times ranked

24

4001 citing authors

#	Article	IF	CITATIONS
1	UV-Associated Mutations Underlie the Etiology of MCV-Negative Merkel Cell Carcinomas. Cancer Research, 2015, 75, 5228-5234.	0.9	270
2	Musket: a multistage <i>k-</i> mer spectrum-based error corrector for Illumina sequence data. Bioinformatics, 2013, 29, 308-315.	4.1	266
3	GRIDSS: sensitive and specific genomic rearrangement detection using positional de Bruijn graph assembly. Genome Research, 2017, 27, 2050-2060.	5.5	255
4	Modelling human blastocysts by reprogramming fibroblasts into iBlastoids. Nature, 2021, 591, 627-632.	27.8	211
5	The Architecture and Evolution of Cancer Neochromosomes. Cancer Cell, 2014, 26, 653-667.	16.8	161
6	Correcting errors in short reads by multiple alignments. Bioinformatics, 2011, 27, 1455-1461.	4.1	145
7	SHREC: a short-read error correction method. Bioinformatics, 2009, 25, 2157-2163.	4.1	133
8	ReprogrammingÂroadmap reveals route toÂhuman induced trophoblast stem cells. Nature, 2020, 586, 101-107.	27.8	131
9	Socrates: identification of genomic rearrangements in tumour genomes by re-aligning soft clipped reads. Bioinformatics, 2014, 30, 1064-1072.	4.1	75
10	Nicotinamide riboside attenuates age-associated metabolic and functional changes in hematopoietic stem cells. Nature Communications, 2021, 12, 2665.	12.8	45
11	Inferring structural variant cancer cell fraction. Nature Communications, 2020, 11, 730.	12.8	33
12	The Scalloped and Nerfin-1 Transcription Factors Cooperate to Maintain Neuronal Cell Fate. Cell Reports, 2018, 25, 1561-1576.e7.	6.4	31
13	HECTOR: a parallel multistage homopolymer spectrum based error corrector for 454 sequencing data. BMC Bioinformatics, 2014, 15, 131.	2.6	25
14	Reference-Free Validation of Short Read Data. PLoS ONE, 2010, 5, e12681.	2.5	23
15	A CX3CR1 Reporter hESC Line Facilitates Integrative Analysis of In-Vitro-Derived Microglia and Improved Microglia Identity upon Neuron-Glia Co-culture. Stem Cell Reports, 2020, 14, 1018-1032.	4.8	16
16	Overview of Fusion Detection Strategies Using Next-Generation Sequencing. Methods in Molecular Biology, 2019, 1908, 125-138.	0.9	15
17	Embryonic Lethality in Homozygous Human Her-2 Transgenic Mice Due to Disruption of the Pds5b Gene. PLoS ONE, 2015, 10, e0136817.	2.5	14
18	Type 2 immune polarization is associated with cardiopulmonary disease in preterm infants. Science Translational Medicine, 2022, 14, eaaz8454.	12.4	14

#	Article	IF	CITATION
19	Expression of a Chimeric Antigen Receptor in Multiple Leukocyte Lineages in Transgenic Mice. PLoS ONE, 2015, 10, e0140543.	2.5	12
20	TINCâ€" A Method to Dissect Regulatory Complexes at Single-Locus Resolutionâ€" Reveals an Extensive Protein Complex at the Nanog Promoter. Stem Cell Reports, 2020, 15, 1246-1259.	4.8	12
21	A Massively Parallel Architecture for Bioinformatics. Lecture Notes in Computer Science, 2009, , 994-1003.	1.3	11
22	Transcriptional signature in microglia isolated from an Alzheimer's disease mouse model treated with scanning ultrasound. Bioengineering and Translational Medicine, 2023, 8, .	7.1	7
23	Improving the Power of Structural Variation Detection by Augmenting the Reference. PLoS ONE, 2015, 10, e0136771.	2.5	6
24	Intestinal stem cell aging signature reveals a reprogramming strategy to enhance regenerative potential. Npj Regenerative Medicine, 2022, 7, .	5.2	4