Mohamed Nadhir Djekidel

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/172662/publications.pdf

Version: 2024-02-01

17 papers 809 citations

759233 12 h-index 17 g-index

20 all docs 20 docs citations

times ranked

20

1028 citing authors

#	Article	IF	CITATIONS
1	eccDNAs are apoptotic products with high innate immunostimulatory activity. Nature, 2021, 599, 308-314.	27.8	121
2	Cell type-specific transcriptional programs in mouse prefrontal cortex during adolescence and addiction. Nature Communications, 2019, 10, 4169.	12.8	100
3	Decoding molecular and cellular heterogeneity of mouse nucleus accumbens. Nature Neuroscience, 2021, 24, 1757-1771.	14.8	87
4	Distinct dynamics and functions of H2AK119ub1 and H3K27me3 in mouse preimplantation embryos. Nature Genetics, 2021, 53, 551-563.	21.4	83
5	Myc and Dnmt1 impede the pluripotent to totipotent state transition in embryonic stem cells. Nature Cell Biology, 2019, 21, 835-844.	10.3	82
6	FIND: difFerential chromatin INteractions Detection using a spatial Poisson process. Genome Research, 2018, 28, 412-422.	5.5	69
7	BL-Hi-C is an efficient and sensitive approach for capturing structural and regulatory chromatin interactions. Nature Communications, 2017, 8, 1622.	12.8	60
8	A transcriptional roadmap for 2C-like–to–pluripotent state transition. Science Advances, 2020, 6, eaay5181.	10.3	44
9	Reprogramming of Chromatin Accessibility in Somatic Cell Nuclear Transfer Is DNA Replication Independent. Cell Reports, 2018, 23, 1939-1947.	6.4	30
10	Alterations of specific chromatin conformation affect ATRA-induced leukemia cell differentiation. Cell Death and Disease, 2018, 9, 200.	6.3	29
11	Acute depletion of CTCF rewires genome-wide chromatin accessibility. Genome Biology, 2021, 22, 244.	8.8	29
12	3CPET: finding co-factor complexes from ChIA-PET data using a hierarchical Dirichlet process. Genome Biology, 2015, 16, 288.	8.8	20
13	Cell type–specific mechanism of Setd1a heterozygosity in schizophrenia pathogenesis. Science Advances, 2022, 8, eabm1077.	10.3	16
14	HiCâ€3DViewer: a new tool to visualize Hi data in 3D space. Quantitative Biology, 2017, 5, 183-190.	0.5	14
15	Developing bioimaging and quantitative methods to study 3D genome. Quantitative Biology, 2016, 4, 129-147.	0.5	9
16	Advances in computational ChIAâ€PET data analysis. Quantitative Biology, 2016, 4, 217-225.	0.5	5
17	In vivo nuclear capture and molecular profiling identifies Gmeb1 as a transcriptional regulator essential for dopamine neuron function. Nature Communications, 2019, 10, 2508.	12.8	3