

Tao Ye

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4020934/publications.pdf>

Version: 2024-02-01

9
papers

720
citations

1307594
7
h-index

1474206
9
g-index

10
all docs

10
docs citations

10
times ranked

2124
citing authors

#	ARTICLE	IF	CITATIONS
1	Waves of sumoylation support transcription dynamics during adipocyte differentiation. <i>Nucleic Acids Research</i> , 2022, 50, 1351-1369.	14.5	8
2	Histone H2Bub1 deubiquitylation is essential for mouse development, but does not regulate global RNA polymerase II transcription. <i>Cell Death and Differentiation</i> , 2021, 28, 2385-2403.	11.2	14
3	The related coactivator complexes SAGA and ATAC control embryonic stem cell self-renewal through acetyltransferase-independent mechanisms. <i>Cell Reports</i> , 2021, 36, 109598.	6.4	9
4	Extensive NEUROG3 occupancy in the human pancreatic endocrine gene regulatory network. <i>Molecular Metabolism</i> , 2021, 53, 101313.	6.5	20
5	Homozygous TAF8 mutation in a patient with intellectual disability results in undetectable TAF8 protein, but preserved RNA polymerase II transcription. <i>Human Molecular Genetics</i> , 2018, 27, 2171-2186.	2.9	22
6	UVB Induces a Genome-Wide Acting Negative Regulatory Mechanism That Operates at the Level of Transcription Initiation in Human Cells. <i>PLoS Genetics</i> , 2014, 10, e1004483.	3.5	34
7	Rfx6 Maintains the Functional Identity of Adult Pancreatic $\hat{1}^2$ Cells. <i>Cell Reports</i> , 2014, 9, 2219-2232.	6.4	114
8	Sumoylation at chromatin governs coordinated repression of a transcriptional program essential for cell growth and proliferation. <i>Genome Research</i> , 2013, 23, 1563-1579.	5.5	121
9	seqMINER: an integrated ChIP-seq data interpretation platform. <i>Nucleic Acids Research</i> , 2011, 39, e35-e35.	14.5	377