

Lingyun Song

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8268858/publications.pdf>

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12
papers

2,480
citations

840776

11
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

6144
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrated chromatin and transcriptomic profiling of patient-derived colon cancer organoids identifies personalized drug targets to overcome oxaliplatin resistance. <i>Genes and Diseases</i> , 2021, 8, 203-214.	3.4	10
2	CommonMind Consortium provides transcriptomic and epigenomic data for Schizophrenia and Bipolar Disorder. <i>Scientific Data</i> , 2019, 6, 180.	5.3	149
3	Open Chromatin Profiling in Adipose Tissue Marks Genomic Regions with Functional Roles in Cardiometabolic Traits. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 2521-2533.	1.8	19
4	Transcriptome and epigenome landscape of human cortical development modeled in organoids. <i>Science</i> , 2018, 362, .	12.6	220
5	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , 2018, 362, .	12.6	516
6	Comprehensive functional genomic resource and integrative model for the human brain. <i>Science</i> , 2018, 362, .	12.6	618
7	Pre-established Chromatin Interactions Mediate the Genomic Response to Glucocorticoids. <i>Cell Systems</i> , 2018, 7, 146-160.e7.	6.2	82
8	Microbiota regulate intestinal epithelial gene expression by suppressing the transcription factor Hepatocyte nuclear factor 4 alpha. <i>Genome Research</i> , 2017, 27, 1195-1206.	5.5	101
9	Direct GR Binding Sites Potentiate Clusters of TF Binding across the Human Genome. <i>Cell</i> , 2016, 166, 1269-1281.e19.	28.9	158
10	Genomic analysis reveals distinct mechanisms and functional classes of SOX10-regulated genes in melanocytes. <i>Human Molecular Genetics</i> , 2015, 24, 5433-5450.	2.9	34
11	Genetic variants and cellular stressors associated with exfoliation syndrome modulate promoter activity of a lncRNA within the <i>LOXL1</i> locus. <i>Human Molecular Genetics</i> , 2015, 24, 6552-6563.	2.9	76
12	DNase-seq: A High-Resolution Technique for Mapping Active Gene Regulatory Elements across the Genome from Mammalian Cells. <i>Cold Spring Harbor Protocols</i> , 2010, 2010, pdb.prot5384.	0.3	497