

Alexias Safi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/660980/publications.pdf>

Version: 2024-02-01

17
papers

1,927
citations

759233

12
h-index

1125743

13
g-index

17
all docs

17
docs citations

17
times ranked

5016
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeted long-read sequencing identifies missing disease-causing variation. American Journal of Human Genetics, 2021, 108, 1436-1449.	6.2	105
2	Integrative QTL analysis of gene expression and chromatin accessibility identifies multi-tissue patterns of genetic regulation. PLoS Genetics, 2020, 16, e1008537.	3.5	35
3	Title is missing!. , 2020, 16, e1008537.		0
4	Title is missing!. , 2020, 16, e1008537.		0
5	Title is missing!. , 2020, 16, e1008537.		0
6	Title is missing!. , 2020, 16, e1008537.		0
7	Open Chromatin Profiling in Adipose Tissue Marks Genomic Regions with Functional Roles in Cardiometabolic Traits. G3: Genes, Genomes, Genetics, 2019, 9, 2521-2533.	1.8	19
8	Comparative Serum Challenges Show Divergent Patterns of Gene Expression and Open Chromatin in Human and Chimpanzee. Genome Biology and Evolution, 2018, 10, 826-839.	2.5	21
9	Tissue- and strain-specific effects of a genotoxic carcinogen 1,3-butadiene on chromatin and transcription. Mammalian Genome, 2018, 29, 153-167.	2.2	21
10	Transcriptome and epigenome landscape of human cortical development modeled in organoids. Science, 2018, 362, .	12.6	220
11	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. Science, 2018, 362, .	12.6	516
12	Comprehensive functional genomic resource and integrative model for the human brain. Science, 2018, 362, .	12.6	618
13	Pre-established Chromatin Interactions Mediate the Genomic Response to Glucocorticoids. Cell Systems, 2018, 7, 146-160.e7.	6.2	82
14	Variation in DNA-Damage Responses to an Inhalational Carcinogen (1,3-Butadiene) in Relation to Strain-Specific Differences in Chromatin Accessibility and Gene Transcription Profiles in C57BL/6J and CAST/Eij Mice. Environmental Health Perspectives, 2017, 125, 107006.	6.0	22
15	Direct GR Binding Sites Potentiate Clusters of TF Binding across the Human Genome. Cell, 2016, 166, 1269-1281.e19.	28.9	158
16	Genomic analysis reveals distinct mechanisms and functional classes of SOX10-regulated genes in melanocytes. Human Molecular Genetics, 2015, 24, 5433-5450.	2.9	34
17	Genetic variants and cellular stressors associated with exfoliation syndrome modulate promoter activity of a lncRNA within the <i>LOXL1</i> locus. Human Molecular Genetics, 2015, 24, 6552-6563.	2.9	76