

Annie W Shieh

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3243197/publications.pdf>

Version: 2024-02-01

16
papers

2,655
citations

1162367

8
h-index

1372195

10
g-index

16
all docs

16
docs citations

16
times ranked

5793
citing authors

#	ARTICLE	IF	CITATIONS
1	Transcriptome-wide isoform-level dysregulation in ASD, schizophrenia, and bipolar disorder. <i>Science</i> , 2018, 362, .	6.0	805
2	Comprehensive functional genomic resource and integrative model for the human brain. <i>Science</i> , 2018, 362, .	6.0	618
3	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , 2018, 362, .	6.0	516
4	The PsychENCODE project. <i>Nature Neuroscience</i> , 2015, 18, 1707-1712.	7.1	371
5	Evaluation of chromatin accessibility in prefrontal cortex of individuals with schizophrenia. <i>Nature Communications</i> , 2018, 9, 3121.	5.8	141
6	The transcription factor POU3F2 regulates a gene coexpression network in brain tissue from patients with psychiatric disorders. <i>Science Translational Medicine</i> , 2018, 10, .	5.8	81
7	A Myogenic Double-Reporter Human Pluripotent Stem Cell Line Allows Prospective Isolation of Skeletal Muscle Progenitors. <i>Cell Reports</i> , 2018, 25, 1966-1981.e4.	2.9	61
8	Neuronal and glial 3D chromatin architecture informs the cellular etiology of brain disorders. <i>Nature Communications</i> , 2021, 12, 3968.	5.8	48
9	DRAMS: A tool to detect and re-align mixed-up samples for integrative studies of multi-omics data. <i>PLoS Computational Biology</i> , 2020, 16, e1007522.	1.5	8
10	Absence of coding somatic single nucleotide variants within well-known candidate genes in late-onset sporadic Alzheimer's Disease based on the analysis of multi-omics data. <i>Neurobiology of Aging</i> , 2021, 108, 207-209.	1.5	6
11	DRAMS: A tool to detect and re-align mixed-up samples for integrative studies of multi-omics data. , 2020, 16, e1007522.		0
12	DRAMS: A tool to detect and re-align mixed-up samples for integrative studies of multi-omics data. , 2020, 16, e1007522.		0
13	DRAMS: A tool to detect and re-align mixed-up samples for integrative studies of multi-omics data. , 2020, 16, e1007522.		0
14	DRAMS: A tool to detect and re-align mixed-up samples for integrative studies of multi-omics data. , 2020, 16, e1007522.		0
15	DRAMS: A tool to detect and re-align mixed-up samples for integrative studies of multi-omics data. , 2020, 16, e1007522.		0
16	DRAMS: A tool to detect and re-align mixed-up samples for integrative studies of multi-omics data. , 2020, 16, e1007522.		0