

Joon-Yong An

List of Publications by Year in descending order

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Version: 2024-02-01

33
papers

5,361
citations

393982

19
h-index

395343

33
g-index

45
all docs

45
docs citations

45
times ranked

9588
citing authors

#	ARTICLE	IF	CITATIONS
1	Large-Scale Exome Sequencing Study Implicates Both Developmental and Functional Changes in the Neurobiology of Autism. <i>Cell</i> , 2020, 180, 568-584.e23.	13.5	1,422
2	Transcriptome-wide isoform-level dysregulation in ASD, schizophrenia, and bipolar disorder. <i>Science</i> , 2018, 362, .	6.0	805
3	Comprehensive functional genomic resource and integrative model for the human brain. <i>Science</i> , 2018, 362, .	6.0	618
4	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , 2018, 362, .	6.0	516
5	An analytical framework for whole-genome sequence association studies and its implications for autism spectrum disorder. <i>Nature Genetics</i> , 2018, 50, 727-736.	9.4	235
6	Genome-wide de novo risk score implicates promoter variation in autism spectrum disorder. <i>Science</i> , 2018, 362, .	6.0	234
7	Transcriptome and epigenome landscape of human cortical development modeled in organoids. <i>Science</i> , 2018, 362, .	6.0	220
8	Opposing Effects on Na V 1.2 Function Underlie Differences Between SCN2A Variants Observed in Individuals With Autism Spectrum Disorder or Infantile Seizures. <i>Biological Psychiatry</i> , 2017, 82, 224-232.	0.7	208
9	Neurodevelopmental and neuropsychiatric disorders represent an interconnected molecular system. <i>Molecular Psychiatry</i> , 2014, 19, 294-301.	4.1	188
10	Defining the diverse spectrum of inversions, complex structural variation, and chromothripsis in the morbid human genome. <i>Genome Biology</i> , 2017, 18, 36.	3.8	159
11	Whole genome sequencing in psychiatric disorders: the WGSPD consortium. <i>Nature Neuroscience</i> , 2017, 20, 1661-1668.	7.1	122
12	Whole-Genome and RNA Sequencing Reveal Variation and Transcriptomic Coordination in the Developing Human Prefrontal Cortex. <i>Cell Reports</i> , 2020, 31, 107489.	2.9	91
13	An integrative analysis of non-coding regulatory DNA variations associated with autism spectrum disorder. <i>Molecular Psychiatry</i> , 2019, 24, 1707-1719.	4.1	59
14	Towards a molecular characterization of autism spectrum disorders: an exome sequencing and systems approach. <i>Translational Psychiatry</i> , 2014, 4, e394-e394.	2.4	57
15	Integrative Multi-Omics Approaches in Cancer Research: From Biological Networks to Clinical Subtypes. <i>Molecules and Cells</i> , 2021, 44, 433-443.	1.0	57
16	Neuronal and glial 3D chromatin architecture informs the cellular etiology of brain disorders. <i>Nature Communications</i> , 2021, 12, 3968.	5.8	48
17	Genetic heterogeneity in autism: From single gene to a pathway perspective. <i>Neuroscience and Biobehavioral Reviews</i> , 2016, 68, 442-453.	2.9	46
18	Revealing the brain's molecular architecture. <i>Science</i> , 2018, 362, 1262-1263.	6.0	45

#	ARTICLE	IF	CITATIONS
19	Genetic architecture of autism spectrum disorder: Lessons from large-scale genomic studies. <i>Neuroscience and Biobehavioral Reviews</i> , 2021, 128, 244-257.	2.9	31
20	Developmental dynamics of voltage-gated sodium channel isoform expression in the human and mouse brain. <i>Genome Medicine</i> , 2021, 13, 135.	3.6	19
21	CRISPR screens identify a novel combination treatment targeting BCL-XL and WNT signaling for KRAS/BRAF-mutated colorectal cancers. <i>Oncogene</i> , 2021, 40, 3287-3302.	2.6	18
22	Ferroptosis-Related Genes in Neurodevelopment and Central Nervous System. <i>Biology</i> , 2021, 10, 35.	1.3	17
23	High-throughput characterization of the role of non-B DNA motifs on promoter function. <i>Cell Genomics</i> , 2022, 2, 100111.	3.0	17
24	Homeostatic plasticity fails at the intersection of autism-gene mutations and a novel class of common genetic modifiers. <i>ELife</i> , 2020, 9, .	2.8	14
25	Large-Scale Exome Sequencing Study Implicates Both Developmental and Functional Changes in the Neurobiology of Autism. <i>SSRN Electronic Journal</i> , 0, , .	0.4	12
26	National human genome projects: an update and an agenda. <i>Epidemiology and Health</i> , 2017, 39, e2017045.	0.8	10
27	ANO1 regulates the maintenance of stemness in glioblastoma stem cells by stabilizing EGFRvIII. <i>Oncogene</i> , 2021, 40, 1490-1502.	2.6	9
28	Non-coding de novo mutations in chromatin interactions are implicated in autism spectrum disorder. <i>Molecular Psychiatry</i> , 2022, 27, 4680-4694.	4.1	9
29	Appreciating the Population-wide Impact of Copy Number Variants on Cognition. <i>Biological Psychiatry</i> , 2017, 82, 78-80.	0.7	4
30	Spatio-Temporal Roles of ASD-Associated Variants in Human Brain Development. <i>Genes</i> , 2020, 11, 535.	1.0	3
31	Kdm3b haploinsufficiency impairs the consolidation of cerebellum-dependent motor memory in mice. <i>Molecular Brain</i> , 2021, 14, 106.	1.3	2
32	Infantile esotropia in a family with <i>TUBB3</i> mutation associated congenital fibrosis of extraocular muscles. <i>Ophthalmic Genetics</i> , 2022, 43, 716-719.	0.5	2
33	Identification of Possible Risk Variants of Familial Strabismus Using Exome Sequencing Analysis. <i>Genes</i> , 2021, 12, 75.	1.0	1