

# Hsiao-Tuan Chao

## List of Publications by Year in descending order

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Version: 2024-02-01

35  
papers

3,847  
citations

257450

24  
h-index

345221

36  
g-index

40  
all docs

40  
docs citations

40  
times ranked

6253  
citing authors

#	ARTICLE	IF	CITATIONS
1	Dysfunction in GABA signalling mediates autism-like stereotypies and Rett syndrome phenotypes. <i>Nature</i> , 2010, 468, 263-269.	27.8	1,042
2	MeCP2 Controls Excitatory Synaptic Strength by Regulating Glutamatergic Synapse Number. <i>Neuron</i> , 2007, 56, 58-65.	8.1	439
3	Loss of MeCP2 in aminergic neurons causes cell-autonomous defects in neurotransmitter synthesis and specific behavioral abnormalities. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 21966-21971.	7.1	240
4	Deletion of <i>Mecp2</i> in <i>Sim1</i> -Expressing Neurons Reveals a Critical Role for MeCP2 in Feeding Behavior, Aggression, and the Response to Stress. <i>Neuron</i> , 2008, 59, 947-958.	8.1	230
5	Distinct domains of complexin I differentially regulate neurotransmitter release. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 949-958.	8.2	198
6	MARRVEL: Integration of Human and Model Organism Genetic Resources to Facilitate Functional Annotation of the Human Genome. <i>American Journal of Human Genetics</i> , 2017, 100, 843-853.	6.2	181
7	A partial loss of function allele of Methyl-CpG-binding protein 2 predicts a human neurodevelopmental syndrome. <i>Human Molecular Genetics</i> , 2008, 17, 1718-1727.	2.9	173
8	Model Organisms Facilitate Rare Disease Diagnosis and Therapeutic Research. <i>Genetics</i> , 2017, 207, 9-27.	2.9	165
9	<i>Math1</i> Is Essential for the Development of Hindbrain Neurons Critical for Perinatal Breathing. <i>Neuron</i> , 2009, 64, 341-354.	8.1	146
10	Dendritic Arborization and Spine Dynamics Are Abnormal in the Mouse Model of MECP2 Duplication Syndrome. <i>Journal of Neuroscience</i> , 2013, 33, 19518-19533.	3.6	123
11	Binding of the complexin N terminus to the SNARE complex potentiates synaptic-vesicle fusion. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 568-575.	8.2	113
12	A Syndromic Neurodevelopmental Disorder Caused by De Novo Variants in <i>EBF3</i> . <i>American Journal of Human Genetics</i> , 2017, 100, 128-137.	6.2	96
13	Transcriptome-directed analysis for Mendelian disease diagnosis overcomes limitations of conventional genomic testing. <i>Journal of Clinical Investigation</i> , 2021, 131, .	8.2	87
14	MeCP2: only 100% will do. <i>Nature Neuroscience</i> , 2012, 15, 176-177.	14.8	59
15	Lysosomal Storage and Albinism Due to Effects of a De Novo <i>CLCN7</i> Variant on Lysosomal Acidification. <i>American Journal of Human Genetics</i> , 2019, 104, 1127-1138.	6.2	59
16	Identification of novel candidate disease genes from de novo exonic copy number variants. <i>Genome Medicine</i> , 2017, 9, 83.	8.2	50
17	De Novo Variants in <i>WDR37</i> Are Associated with Epilepsy, Colobomas, Dysmorphism, Developmental Delay, Intellectual Disability, and Cerebellar Hypoplasia. <i>American Journal of Human Genetics</i> , 2019, 105, 413-424.	6.2	43
18	Partial Loss of <i>USP9X</i> Function Leads to a Male Neurodevelopmental and Behavioral Disorder Converging on Transforming Growth Factor $\beta^2$ Signaling. <i>Biological Psychiatry</i> , 2020, 87, 100-112.	1.3	42

#	ARTICLE	IF	CITATIONS
19	Stxbp1/Munc18-1 haploinsufficiency impairs inhibition and mediates key neurological features of STXBP1 encephalopathy. <i>ELife</i> , 2020, 9, .	6.0	42
20	De novo EIF2AK1 and EIF2AK2 Variants Are Associated with Developmental Delay, Leukoencephalopathy, and Neurologic Decompensation. <i>American Journal of Human Genetics</i> , 2020, 106, 570-583.	6.2	37
21	Clinical and molecular characterization of de novo loss of function variants in <i>HNRNP1</i> . <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2680-2689.	1.2	34
22	In Vivo Functional Study of Disease-associated Rare Human Variants Using <i>Drosophila</i> . <i>Journal of Visualized Experiments</i> , 2019, , .	0.3	34
23	The yin and yang of MeCP2 phosphorylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 4577-4578.	7.1	31
24	De Novo Pathogenic Variants in N-cadherin Cause a Syndromic Neurodevelopmental Disorder with Corpus Callosum, Axon, Cardiac, Ocular, and Genital Defects. <i>American Journal of Human Genetics</i> , 2019, 105, 854-868.	6.2	29
25	<i>Drosophila</i> functional screening of de novo variants in autism uncovers damaging variants and facilitates discovery of rare neurodevelopmental diseases. <i>Cell Reports</i> , 2022, 38, 110517.	6.4	24
26	De Novo Variants in CDK19 Are Associated with a Syndrome Involving Intellectual Disability and Epileptic Encephalopathy. <i>American Journal of Human Genetics</i> , 2020, 106, 717-725.	6.2	23
27	Investigation of Synapse Formation and Function in a Glutamatergic-GABAergic Two-Neuron Microcircuit. <i>Journal of Neuroscience</i> , 2014, 34, 855-868.	3.6	22
28	DYRK1A-related intellectual disability: a syndrome associated with congenital anomalies of the kidney and urinary tract. <i>Genetics in Medicine</i> , 2019, 21, 2755-2764.	2.4	19
29	Building dialogues between clinical and biomedical research through cross-species collaborations. <i>Seminars in Cell and Developmental Biology</i> , 2017, 70, 49-57.	5.0	16
30	Heterozygous loss-of-function variants significantly expand the phenotypes associated with loss of GDF11. <i>Genetics in Medicine</i> , 2021, 23, 1889-1900.	2.4	13
31	Recurrent mosaic MTOR c.5930C > T (p.Thr1977Ile) variant causing megalencephaly, asymmetric polymicrogyria, and cutaneous pigmentary mosaicism: Case report and review of the literature. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 475-479.	1.2	11
32	<i>EIF2AK2</i> -related Neurodevelopmental Disorder With Leukoencephalopathy, Developmental Delay, and Episodic Neurologic Regression Mimics Pelizaeus-Merzbacher Disease. <i>Neurology: Genetics</i> , 2021, 7, e539.	1.9	9
33	Magnetic Resonance Imaging characteristics in case of TOR1AIP1 muscular dystrophy. <i>Clinical Imaging</i> , 2019, 58, 108-113.	1.5	6
34	An Integrated Phenotypic and Genotypic Approach Reveals a High-Risk Subtype Association for <i>EBF3</i> Missense Variants Affecting the Zinc Finger Domain. <i>Annals of Neurology</i> , 2022, 92, 138-153.	5.3	5
35	<i>PRUNE1</i> c.933G > A synonymous variant induces exon 7 skipping, disrupts the <i>DHHA2</i> domain, and leads to an atypical <i>NMIHBA</i> syndrome presentation: Case report and review of the literature. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 1868-1874.	1.2	2