

# Ying Zhu

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

26

papers

4,862

citations

19

h-index

27

g-index

27

ext. papers

6,506

ext. citations

22.4

avg, IF

4.44

L-index

#	Paper	IF	Citations
26	Transcriptional programs regulating neuronal differentiation are disrupted in DLG2 knockout human embryonic stem cells and enriched for schizophrenia and related disorders risk variants.. <i>Nature Communications</i> , <b>2022</b> , 13, 27	17.4	3
25	Rapid Body-Wide Transcriptomic Turnover During Rhesus Macaque Perinatal Development. <i>Frontiers in Physiology</i> , <b>2021</b> , 12, 690540	4.6	1
24	TLR9 Deficiency in B Cells Promotes Immune Tolerance via Interleukin-10 in a Type 1 Diabetes Mouse Model. <i>Diabetes</i> , <b>2021</b> , 70, 504-515	0.9	1
23	Differential immunomodulatory effect of PARP inhibition in BRCA1 deficient and competent tumor cells. <i>Biochemical Pharmacology</i> , <b>2021</b> , 184, 114359	6	2
22	Disruption of TCF4 regulatory networks leads to abnormal cortical development and mental disabilities. <i>Molecular Psychiatry</i> , <b>2019</b> , 24, 1235-1246	15.1	34
21	B Cell Presentation of Chlamydia Antigen Selects Out Protective CD4 <sup>hi</sup> T Cells: Implications for Genital Tract Tissue-Resident Memory Lymphocyte Clusters. <i>Infection and Immunity</i> , <b>2018</b> , 86,	3.7	20
20	Transcriptome and epigenome landscape of human cortical development modeled in organoids. <i>Science</i> , <b>2018</b> , 362,	33.3	142
19	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , <b>2018</b> , 362,	33.3	277
18	Spatiotemporal transcriptomic divergence across human and macaque brain development. <i>Science</i> , <b>2018</b> , 362,	33.3	127
17	Transcriptome-wide isoform-level dysregulation in ASD, schizophrenia, and bipolar disorder. <i>Science</i> , <b>2018</b> , 362,	33.3	434
16	Comprehensive functional genomic resource and integrative model for the human brain. <i>Science</i> , <b>2018</b> , 362,	33.3	319
15	Molecular and cellular reorganization of neural circuits in the human lineage. <i>Science</i> , <b>2017</b> , 358, 1027-1033,	33.3	127
14	Zika Virus Disrupts Phospho-TBK1 Localization and Mitosis in Human Neuroepithelial Stem Cells and Radial Glia. <i>Cell Reports</i> , <b>2016</b> , 16, 2576-2592	10.6	192
13	The Molecular Landscape of the Developing Human Central Nervous System <b>2016</b> , 203-220		1
12	The Cellular and Molecular Landscapes of the Developing Human Central Nervous System. <i>Neuron</i> , <b>2016</b> , 89, 248-68	13.9	312
11	Down Syndrome Developmental Brain Transcriptome Reveals Defective Oligodendrocyte Differentiation and Myelination. <i>Neuron</i> , <b>2016</b> , 89, 1208-1222	13.9	120
10	Simultaneous dimension reduction and adjustment for confounding variation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2016</b> , 113, 14662-14667	11.5	24

9	TSHZ3 deletion causes an autism syndrome and defects in cortical projection neurons. <i>Nature Genetics</i> , <b>2016</b> , 48, 1359-1369	36.3	36
8	Temporal specification and bilaterality of human neocortical topographic gene expression. <i>Neuron</i> , <b>2014</b> , 81, 321-32	13.9	159
7	XSAAnno: a framework for building ortholog models in cross-species transcriptome comparisons. <i>BMC Genomics</i> , <b>2014</b> , 15, 343	4.5	17
6	Coexpression networks implicate human midfetal deep cortical projection neurons in the pathogenesis of autism. <i>Cell</i> , <b>2013</b> , 155, 997-1007	56.2	591
5	Recessive LAMC3 mutations cause malformations of occipital cortical development. <i>Nature Genetics</i> , <b>2011</b> , 43, 590-4	36.3	85
4	Spatio-temporal transcriptome of the human brain. <i>Nature</i> , <b>2011</b> , 478, 483-9	50.4	1302
3	TBR1 directly represses Fezf2 to control the laminar origin and development of the corticospinal tract. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2011</b> , 108, 3041-6	11.5	140
2	Whole-exome sequencing identifies recessive WDR62 mutations in severe brain malformations. <i>Nature</i> , <b>2010</b> , 467, 207-10	50.4	395
1	DLG2 knockout reveals neurogenic transcriptional programs underlying neuropsychiatric disorders and cognition		1