

# Amanda J Price

## 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.

18  
papers

1,595  
citations

10  
h-index

18  
g-index

18  
ext. papers

2,456  
ext. citations

19.1  
avg. IF

3.04  
L-index

#	Paper	IF	Citations
18	Cortical cellular diversity and development in schizophrenia. <i>Molecular Psychiatry</i> , <b>2021</b> , 26, 203-217	15.1	3
17	Characterizing the dynamic and functional DNA methylation landscape in the developing human cortex. <i>Epigenetics</i> , <b>2021</b> , 16, 1-13	5.7	9
16	Genome-wide sequencing-based identification of methylation quantitative trait loci and their role in schizophrenia risk. <i>Nature Communications</i> , <b>2021</b> , 12, 5251	17.4	3
15	Dissecting transcriptomic signatures of neuronal differentiation and maturation using iPSCs. <i>Nature Communications</i> , <b>2020</b> , 11, 462	17.4	37
14	Characterizing the nuclear and cytoplasmic transcriptomes in developing and mature human cortex uncovers new insight into psychiatric disease gene regulation. <i>Genome Research</i> , <b>2020</b> , 30, 1-11	9.7	12
13	Schizophrenia risk variants influence multiple classes of transcripts of sorting nexin 19 (SNX19). <i>Molecular Psychiatry</i> , <b>2020</b> , 25, 831-843	15.1	17
12	Divergent neuronal DNA methylation patterns across human cortical development reveal critical periods and a unique role of CpH methylation. <i>Genome Biology</i> , <b>2019</b> , 20, 196	18.3	42
11	Regional Heterogeneity in Gene Expression, Regulation, and Coherence in the Frontal Cortex and Hippocampus across Development and Schizophrenia. <i>Neuron</i> , <b>2019</b> , 103, 203-216.e8	13.9	67
10	Transcriptome and epigenome landscape of human cortical development modeled in organoids. <i>Science</i> , <b>2018</b> , 362,	33.3	142
9	Integrative functional genomic analysis of human brain development and neuropsychiatric risks. <i>Science</i> , <b>2018</b> , 362,	33.3	277
8	Transcriptome-wide isoform-level dysregulation in ASD, schizophrenia, and bipolar disorder. <i>Science</i> , <b>2018</b> , 362,	33.3	434
7	Comprehensive functional genomic resource and integrative model for the human brain. <i>Science</i> , <b>2018</b> , 362,	33.3	319
6	Engineering targeted chromosomal amplifications in human breast epithelial cells. <i>Breast Cancer Research and Treatment</i> , <b>2015</b> , 152, 313-21	4.4	2
5	The PsychENCODE project. <i>Nature Neuroscience</i> , <b>2015</b> , 18, 1707-12	25.5	226
4	Widespread methylation quantitative trait loci and their role in schizophrenia risk		1
3	Divergent neuronal DNA methylation patterns across human cortical development: Critical periods and a unique role of CpH methylation		1
2	Characterizing the nuclear and cytoplasmic transcriptomes in developing and mature human cortex uncovers new insight into psychiatric disease gene regulation		2

1 Characterizing the dynamic and functional DNA methylation landscape in the developing human cortex 1