

# Jie Huang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9020781/publications.pdf>

Version: 2024-02-01

15  
papers

2,029  
citations

758635

12  
h-index

996533

15  
g-index

16  
all docs

16  
docs citations

16  
times ranked

4313  
citing authors

#	ARTICLE	IF	CITATIONS
1	Direct Conversion of Fibroblasts to Neurons by Reprogramming PTB-Regulated MicroRNA Circuits. <i>Cell</i> , 2013, 152, 82-96.	13.5	508
2	MicroRNA Directly Enhances Mitochondrial Translation during Muscle Differentiation. <i>Cell</i> , 2014, 158, 607-619.	13.5	385
3	SR Proteins Collaborate with 7SK and Promoter-Associated Nascent RNA to Release Paused Polymerase. <i>Cell</i> , 2013, 153, 855-868.	13.5	279
4	Genome-wide Analysis Reveals SR Protein Cooperation and Competition in Regulated Splicing. <i>Molecular Cell</i> , 2013, 50, 223-235.	4.5	261
5	Mechanisms for U2AF to define 3' splice sites and regulate alternative splicing in the human genome. <i>Nature Structural and Molecular Biology</i> , 2014, 21, 997-1005.	3.6	150
6	Nuclear Matrix Factor hnRNP U/SAF-A Exerts a Global Control of Alternative Splicing by Regulating U2 snRNP Maturation. <i>Molecular Cell</i> , 2012, 45, 656-668.	4.5	146
7	Genome-wide analyses of chromatin interactions after the loss of Pol I, Pol II, and Pol III. <i>Genome Biology</i> , 2020, 21, 158.	3.8	89
8	Deep Insight into the <i>Ganoderma lucidum</i> by Comprehensive Analysis of Its Transcriptome. <i>PLoS ONE</i> , 2012, 7, e44031.	1.1	60
9	Transcriptome and Proteome Exploration to Provide a Resource for the Study of <i>Agrocybe aegerita</i> . <i>PLoS ONE</i> , 2013, 8, e56686.	1.1	56
10	The RNA binding protein EWS is broadly involved in the regulation of pri-miRNA processing in mammalian cells. <i>Nucleic Acids Research</i> , 2017, 45, 12481-12495.	6.5	26
11	Acetylglutamate kinase is required for both gametophyte function and embryo development in <i>Arabidopsis thaliana</i> . <i>Journal of Integrative Plant Biology</i> , 2017, 59, 642-656.	4.1	20
12	A structured RNA in hepatitis B virus post-transcriptional regulatory element represses alternative splicing in a sequence-independent and position-dependent manner. <i>FEBS Journal</i> , 2011, 278, 1533-1546.	2.2	16
13	BAT Hi-C maps global chromatin interactions in an efficient and economical way. <i>Methods</i> , 2020, 170, 38-47.	1.9	13
14	Multiplex Analysis of PolyA-Linked Sequences (MAPS): An RNA-Seq Strategy to Profile Poly(A+) RNA. <i>Methods in Molecular Biology</i> , 2014, 1125, 169-178.	0.4	10
15	BRD2 interconnects with BRD3 to facilitate Pol II transcription initiation and elongation to prime promoters for cell differentiation. <i>Cellular and Molecular Life Sciences</i> , 2022, 79, .	2.4	9