

# Yajing Hao

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7043023/publications.pdf>

Version: 2024-02-01

20  
papers

1,777  
citations

567281

15  
h-index

713466

21  
g-index

24  
all docs

24  
docs citations

24  
times ranked

3210  
citing authors

#	ARTICLE	IF	CITATIONS
1	NONCODE 2016: an informative and valuable data source of long non-coding RNAs. <i>Nucleic Acids Research</i> , 2016, 44, D203-D208.	14.5	574
2	Single-cell transcriptomic architecture and intercellular crosstalk of human intrahepatic cholangiocarcinoma. <i>Journal of Hepatology</i> , 2020, 73, 1118-1130.	3.7	272
3	Mesenchymal Stem Cells Promote Hepatocarcinogenesis via lncRNA-MUF Interaction with ANXA2 and miR-34a. <i>Cancer Research</i> , 2017, 77, 6704-6716.	0.9	193
4	NPInter v3.0: an upgraded database of noncoding RNA-associated interactions. <i>Database: the Journal of Biological Databases and Curation</i> , 2016, 2016, baw057.	3.0	130
5	NPInter v4.0: an integrated database of ncRNA interactions. <i>Nucleic Acids Research</i> , 2020, 48, D160-D165.	14.5	106
6	The long non-coding RNA lncHDAC2 drives the self-renewal of liver cancer stem cells via activation of Hedgehog signaling. <i>Journal of Hepatology</i> , 2019, 70, 918-929.	3.7	93
7	SmProt: a database of small proteins encoded by annotated coding and non-coding RNA loci. <i>Briefings in Bioinformatics</i> , 2018, 19, bbx005.	6.5	85
8	Long noncoding RNA LINC01186, regulated by TGF- $\beta$ /SMAD3, inhibits migration and invasion through Epithelial-Mesenchymal-Transition in lung cancer. <i>Gene</i> , 2017, 608, 1-12.	2.2	50
9	TGF- $\beta$ /SMAD4-Regulated lncRNA-LINP1 Inhibits Epithelial-Mesenchymal Transition in Lung Cancer. <i>International Journal of Biological Sciences</i> , 2018, 14, 1715-1723.	6.4	43
10	MicroRNA-379-5p is associated with biochemical premature ovarian insufficiency through PARP1 and XRCC6. <i>Cell Death and Disease</i> , 2018, 9, 106.	6.3	42
11	lncVar: a database of genetic variation associated with long non-coding genes. <i>Bioinformatics</i> , 2017, 33, 112-118.	4.1	33
12	lincK contributes to breast tumorigenesis by promoting proliferation and epithelial-to-mesenchymal transition. <i>Journal of Hematology and Oncology</i> , 2019, 12, 19.	17.0	29
13	SmProt: A Reliable Repository with Comprehensive Annotation of Small Proteins Identified from Ribosome Profiling. <i>Genomics, Proteomics and Bioinformatics</i> , 2021, 19, 602-610.	6.9	28
14	Profiling the long noncoding RNA interaction network in the regulatory elements of target genes by chromatin in situ reverse transcription sequencing. <i>Genome Research</i> , 2019, 29, 1521-1532.	5.5	27
15	Deeply Mining a Universe of Peptides Encoded by Long Noncoding RNAs. <i>Molecular and Cellular Proteomics</i> , 2021, 20, 100109.	3.8	24
16	The Landscape of Viral Expression Reveals Clinically Relevant Viruses with Potential Capability of Promoting Malignancy in Lower-Grade Glioma. <i>Clinical Cancer Research</i> , 2017, 23, 2177-2185.	7.0	12
17	Chromatin lncRNA Platr10 controls stem cell pluripotency by coordinating an intrachromosomal regulatory network. <i>Genome Biology</i> , 2021, 22, 233.	8.8	12
18	Active retrotransposons help maintain pericentromeric heterochromatin required for faithful cell division. <i>Genome Research</i> , 2020, 30, 1570-1582.	5.5	9

#	ARTICLE	IF	CITATIONS
19	Differential long non-coding RNA and mRNA expression in differentiated human glioblastoma stem cells. <i>Molecular Medicine Reports</i> , 2016, 14, 2067-2076.	2.4	7
20	LncVar: Deciphering Genetic Variations Associated with Long Noncoding Genes. <i>Methods in Molecular Biology</i> , 2019, 1870, 189-198.	0.9	3