

# Jie He

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2261082/publications.pdf>

Version: 2024-02-01

13  
papers

161  
citations

1163117

8  
h-index

1281871

11  
g-index

15  
all docs

15  
docs citations

15  
times ranked

182  
citing authors

#	ARTICLE	IF	CITATIONS
1	High expression of long noncoding RNA HULC is a poor predictor of prognosis and regulates cell proliferation in glioma. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 113-120.	2.0	30
2	Screening of Nanobody Specific for Peanut Major Allergen Ara h 3 by Phage Display. <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 11219-11229.	5.2	20
3	Tumor-associated macrophages of the M1/M2 phenotype are involved in the regulation of malignant biological behavior of breast cancer cells through the EMT pathway. <i>Medical Oncology</i> , 2022, 39, 83.	2.5	19
4	Immunomagnetic microscopy of tumor tissues using quantum sensors in diamond. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	17
5	Over expression of PTEN induces apoptosis and prevents cell proliferation in breast cancer cells. <i>Acta Biochimica Polonica</i> , 2020, 67, 515-519.	0.5	16
6	Knockdown of Ubiquitin Associated Protein 2-Like (UBAP2L) Inhibits Growth and Metastasis of Hepatocellular Carcinoma. <i>Medical Science Monitor</i> , 2018, 24, 7109-7118.	1.1	12
7	Long non-coding RNA HULC stimulates the epithelial-mesenchymal transition process and vasculogenic mimicry in human glioblastoma. <i>Cancer Medicine</i> , 2021, 10, 5270-5282.	2.8	11
8	Clinicopathological characteristics and genetic variations of uterine tumours resembling ovarian sex cord tumours. <i>Journal of Clinical Pathology</i> , 2022, 75, 776-781.	2.0	10
9	Nuclei R-CNN: Improve Mask R-CNN for Nuclei Segmentation. , 2019, , .		8
10	&lt;p&gt;Quantitative Proteomics Analysis Indicates That Upregulation of lncRNA HULC Promotes Pathogenesis of Glioblastoma Cells&lt;/p&gt;. <i>OncoTargets and Therapy</i> , 2020, Volume 13, 5927-5938.	2.0	6
11	Quantitative proteomics analysis of glioblastoma cell lines after lncRNA HULC silencing. <i>Scientific Reports</i> , 2021, 11, 12587.	3.3	6
12	LSCDFS-MKL: A multiple kernel based method for lung squamous cell carcinomas disease-free survival prediction with pathological and genomic data. <i>Journal of Biomedical Informatics</i> , 2019, 94, 103194.	4.3	5
13	Aberrant activation of Wnt/catenin signaling and overexpression of ABCG2 contributes to apoptosis down regulation and tumor progression of high grade ovarian cancer. <i>Acta Biochimica Polonica</i> , 2021, 68, 127-133.	0.5	0