

# Yao Qianlan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7533777/publications.pdf>

Version: 2024-02-01

8  
papers

90  
citations

1937685

4  
h-index

1720034

7  
g-index

8  
all docs

8  
docs citations

8  
times ranked

177  
citing authors

#	ARTICLE	IF	CITATIONS
1	Kinaseâ€‘substrate Edge Biomarkers Provide A More Accurate Prognostic Prediction in ER-negative Breast Cancer. <i>Genomics, Proteomics and Bioinformatics</i> , 2020, 18, 525-538.	6.9	0
2	Systematic identification of rabbit LncRNAs reveals functional roles in atherosclerosis. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2018, 1864, 2266-2273.	3.8	2
3	Comparative transcriptomics reveals specific responding genes associated with atherosclerosis in rabbit and mouse models. <i>PLoS ONE</i> , 2018, 13, e0201618.	2.5	3
4	Global Prioritizing Disease Candidate lncRNAs via a Multi-level Composite Network. <i>Scientific Reports</i> , 2017, 7, 39516.	3.3	47
5	Protein coding gene CRNKL1 as a potential prognostic biomarker in esophageal adenocarcinoma. <i>Artificial Intelligence in Medicine</i> , 2017, 76, 1-6.	6.5	4
6	Population-based study of effectiveness of neoadjuvant radiotherapy on survival in US rectal cancer patients according to age. <i>Scientific Reports</i> , 2017, 7, 3471.	3.3	8
7	Comprehensive analysis of differential co-expression patterns reveal transcriptional dysregulation mechanism and identify novel prognostic lncRNAs in esophageal squamous cell carcinoma. <i>OncoTargets and Therapy</i> , 2017, Volume 10, 3095-3105.	2.0	22
8	RWCFusion: identifying phenotype-specific cancer driver gene fusions based on fusion pair random walk scoring method. <i>Oncotarget</i> , 2016, 7, 61054-61068.	1.8	4