

# Siyu Han

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1777518/publications.pdf>

Version: 2024-02-01

11  
papers

1,514  
citations

1039880

9  
h-index

1372474

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

3779  
citing authors

#	ARTICLE	IF	CITATIONS
1	Long Noncoding RNA Identification: Comparing Machine Learning Based Tools for Long Noncoding Transcripts Discrimination. <i>BioMed Research International</i> , 2016, 2016, 1-14.	0.9	1,176
2	LncFinder: an integrated platform for long non-coding RNA identification utilizing sequence intrinsic composition, structural information and physicochemical property. <i>Briefings in Bioinformatics</i> , 2019, 20, 2009-2027.	3.2	98
3	RPITER: A Hierarchical Deep Learning Framework for ncRNA-Protein Interaction Prediction. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1070.	1.8	55
4	LncIdent: A Tool for Rapid Identification of Long Noncoding RNAs Utilizing Sequence Intrinsic Composition and Open Reading Frame Information. <i>International Journal of Genomics</i> , 2016, 2016, 1-11.	0.8	43
5	Predictive biomarkers of colorectal cancer. <i>Computational Biology and Chemistry</i> , 2019, 83, 107106.	1.1	34
6	Classifying Breast Cancer Subtypes Using Multiple Kernel Learning Based on Omics Data. <i>Genes</i> , 2019, 10, 200.	1.0	34
7	Long Noncoding RNA and Protein Interactions: From Experimental Results to Computational Models Based on Network Methods. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1284.	1.8	29
8	Capsule-LPI: a LncRNA-protein interaction predicting tool based on a capsule network. <i>BMC Bioinformatics</i> , 2021, 22, 246.	1.2	26
9	Predicting lncRNA-disease associations using network topological similarity based on deep mining heterogeneous networks. <i>Mathematical Biosciences</i> , 2019, 315, 108229.	0.9	17
10	LPIInsider: a webserver for lncRNA-protein interaction extraction from the literature. <i>BMC Bioinformatics</i> , 2022, 23, 135.	1.2	2
11	Online Genomic Resources and Bioinformatics Tools Available for Epigenetics and Non-coding RNA. , 2021, , 306-328.		0