

Jukun Song

List of Publications by Citations

Source: <https://exaly.com/author-pdf/8398596/jukun-song-publications-by-citations.pdf>

Version: 2024-04-25

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

22

papers

318

citations

8

h-index

17

g-index

26

ext. papers

494

ext. citations

5.1

avg. IF

3.71

L-index

#	Paper	IF	Citations
22	Reconstruction and analysis of the aberrant lncRNA-miRNA-mRNA network based on competitive endogenous RNA in CESC. <i>Journal of Cellular Biochemistry</i> , 2018 , 119, 6665-6673	4.7	62
21	Systematic analysis of ACE2 and TMPRSS2 expression in salivary glands reveals underlying transmission mechanism caused by SARS-CoV-2. <i>Journal of Medical Virology</i> , 2020 , 92, 2556-2566	19.7	47
20	Five key lncRNAs considered as prognostic targets for predicting pancreatic ductal adenocarcinoma. <i>Journal of Cellular Biochemistry</i> , 2018 , 119, 4559-4569	4.7	41
19	Systematic analysis of alternative splicing signature unveils prognostic predictor for kidney renal clear cell carcinoma. <i>Journal of Cellular Physiology</i> , 2019 , 234, 22753-22764	7	29
18	The relationship between tooth loss and mortality from all causes, cardiovascular diseases, and coronary heart disease in the general population: systematic review and dose-response meta-analysis of prospective cohort studies. <i>Bioscience Reports</i> , 2019 , 39,	4.1	23
17	Patterns of Immune Infiltration in HNC and Their Clinical Implications: A Gene Expression-Based Study. <i>Frontiers in Oncology</i> , 2019 , 9, 1285	5.3	20
16	Systematic analysis of survival-associated alternative splicing signatures uncovers prognostic predictors for head and neck cancer. <i>Journal of Cellular Physiology</i> , 2019 , 234, 15836	7	16
15	Systematic analysis of differentially methylated expressed genes and site-specific methylation as potential prognostic markers in head and neck cancer. <i>Journal of Cellular Physiology</i> , 2019 , 234, 22687-22702	7	12
14	Identification of three molecular subtypes based on immune infiltration in ovarian cancer and its prognostic value. <i>Bioscience Reports</i> , 2020 , 40,	4.1	8
13	Expression and Role of Dickkopf-1 (Dkk1) in Tumors: From the Cells to the Patients. <i>Cancer Management and Research</i> , 2021 , 13, 659-675	3.6	8
12	Development and validation of an immune-related gene prognostic model for stomach adenocarcinoma. <i>Bioscience Reports</i> , 2020 , 40,	4.1	7
11	Combined Analysis of the Aberrant Epigenetic Alteration of Pancreatic Ductal Adenocarcinoma. <i>BioMed Research International</i> , 2019 , 2019, 9379864	3	7
10	Genomic analysis and clinical implications of immune cell infiltration in gastric cancer. <i>Bioscience Reports</i> , 2020 , 40,	4.1	6
9	Effects of CPAP on the transcriptional signatures in patients with obstructive sleep apnea via coexpression network analysis. <i>Journal of Cellular Biochemistry</i> , 2019 , 120, 9277-9290	4.7	5
8	Co-expression of key gene modules and pathways of human breast cancer cell lines. <i>Bioscience Reports</i> , 2019 , 39,	4.1	5
7	miRNA-Based Feature Classifier Is Associated with Tumor Mutational Burden in Head and Neck Squamous Cell Carcinoma. <i>BioMed Research International</i> , 2020 , 2020, 1686480	3	5
6	Immunogenomic Profiling and Classification of Prostate Cancer Based on HIF-1 Signaling Pathway. <i>Frontiers in Oncology</i> , 2020 , 10, 1374	5.3	5

5	Association between tooth loss and risk of oesophageal cancer: a dose-response meta-analysis. <i>SpringerPlus</i> , 2016 , 5, 1020		4
4	Extracellular matrix-related genes play an important role in the progression of NMIBC to MIBC: a bioinformatics analysis study. <i>Bioscience Reports</i> , 2020 , 40,	4.1	4
3	Identification of immune-based prostate cancer subtypes using mRNA expression. <i>Bioscience Reports</i> , 2021 , 41,	4.1	2
2	Systematic analysis of the function and prognostic value of RNA binding protein in head and neck squamous cell carcinoma. <i>European Archives of Oto-Rhino-Laryngology</i> , 2021 , 1	3.5	1
1	Classification and clinical value of three immune subtypes of ovarian cancer based on transcriptome data. <i>International Journal of Transgender Health</i> , 2021 , 14, 963-975	3	0