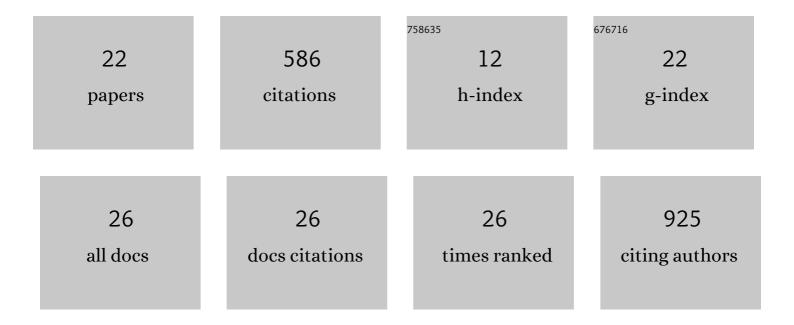
Jukun Song

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Systematic analysis of ACE2 and TMPRSS2 expression in salivary glands reveals underlying transmission mechanism caused by SARSâ€CoVâ€2. Journal of Medical Virology, 2020, 92, 2556-2566.	2.5	86
2	Reconstruction and analysis of the aberrant lncRNAâ€miRNAâ€mRNA network based on competitive endogenous RNA in CESC. Journal of Cellular Biochemistry, 2018, 119, 6665-6673.	1.2	82
3	Five key IncRNAs considered as prognostic targets for predicting pancreatic ductal adenocarcinoma. Journal of Cellular Biochemistry, 2018, 119, 4559-4569.	1.2	65
4	Systematic analysis of alternative splicing signature unveils prognostic predictor for kidney renal clear cell carcinoma. Journal of Cellular Physiology, 2019, 234, 22753-22764.	2.0	58
5	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. Bioscience Reports, 2019, 39, .	1.1	55
6	Expression and Role of Dickkopf-1 (Dkk1) in Tumors: From the Cells to the Patients. Cancer Management and Research, 2021, Volume 13, 659-675.	0.9	34
7	Patterns of Immune Infiltration in HNC and Their Clinical Implications: A Gene Expression-Based Study. Frontiers in Oncology, 2019, 9, 1285.	1.3	31
8	Systematic analysis of survivalâ€associated alternative splicing signatures uncovers prognostic predictors for head and neck cancer. Journal of Cellular Physiology, 2019, 234, 15836-15846.	2.0	26
9	Systematic analysis of differentially methylated expressed genes and siteâ€speciff methylation as potential prognostic markers in head and neck cancer. Journal of Cellular Physiology, 2019, 234, 22687-22702.	2.0	22
10	Development and validation of an immune-related gene prognostic model for stomach adenocarcinoma. Bioscience Reports, 2020, 40, .	1.1	17
11	Identification of three molecular subtypes based on immune infiltration in ovarian cancer and its prognostic value. Bioscience Reports, 2020, 40, .	1.1	17
12	Co-expression of key gene modules and pathways of human breast cancer cell lines. Bioscience Reports, 2019, 39, .	1.1	13
13	Combined Analysis of the Aberrant Epigenetic Alteration of Pancreatic Ductal Adenocarcinoma. BioMed Research International, 2019, 2019, 1-11.	0.9	13
14	Extracellular matrix-related genes play an important role in the progression of NMIBC to MIBC: a bioinformatics analysis study. Bioscience Reports, 2020, 40, .	1.1	13
15	Genomic analysis and clinical implications of immune cell infiltration in gastric cancer. Bioscience Reports, 2020, 40, .	1.1	12
16	Immunogenomic Profiling and Classification of Prostate Cancer Based on HIF-1 Signaling Pathway. Frontiers in Oncology, 2020, 10, 1374.	1.3	9
17	miRNA-Based Feature Classifier Is Associated with Tumor Mutational Burden in Head and Neck Squamous Cell Carcinoma. BioMed Research International, 2020, 2020, 1-14.	0.9	8
18	Identification of immune-based prostate cancer subtypes using mRNA expression. Bioscience Reports, 2021, 41, .	1.1	7

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#	Article	IF	CITATIONS
19	Effects of CPAP on the transcriptional signatures in patients with obstructive sleep apnea via coexpression network analysis. Journal of Cellular Biochemistry, 2019, 120, 9277-9290.	1.2	6
20	Association between tooth loss and risk of oesophageal cancer: a dose–response meta-analysis. SpringerPlus, 2016, 5, 1020.	1.2	4
21	Systematic analysis of the function and prognostic value of RNA binding protein in head and neck squamous cell carcinoma. European Archives of Oto-Rhino-Laryngology, 2021, , 1.	0.8	2
22	Classification and clinical value of three immune subtypes of ovarian cancer based on transcriptome data. International Journal of Transgender Health, 2021, 14, 963-975.	1.1	2