Yan Sun

List of Publications by Year in Descending Order

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

34	1,047	15	32
papers	citations	h-index	g-index
36 ext. papers	1,328 ext. citations	6.6 avg, IF	4·47 L-index

#	Paper	IF	Citations
34	Rack1 regulates pro-inflammatory cytokines by NF- B in diabetic nephropathy. <i>Open Medicine</i> (<i>Poland</i>), 2022 , 17, 978-990	2.2	O
33	A novel lncRNA ROPM-mediated lipid metabolism governs breast cancer stem cell properties. Journal of Hematology and Oncology, 2021 , 14, 178	22.4	10
32	Identification of potential oncogenes in triple-negative breast cancer based on bioinformatics analyses. <i>Oncology Letters</i> , 2021 , 21, 363	2.6	O
31	RNA-Seq analysis reveals critical transcriptome changes caused by sodium butyrate in DN mouse models. <i>Bioscience Reports</i> , 2021 , 41,	4.1	2
30	Systematic Identification of Survival-Associated Alternative Splicing Events in Kidney Renal Clear Cell Carcinoma. <i>Computational and Mathematical Methods in Medicine</i> , 2021 , 2021, 5576933	2.8	1
29	SLC6A8-mediated intracellular creatine accumulation enhances hypoxic breast cancer cell survival via ameliorating oxidative stress. <i>Journal of Experimental and Clinical Cancer Research</i> , 2021 , 40, 168	12.8	2
28	Identification of ribosomal protein family in triple-negative breast cancer by bioinformatics analysis. <i>Bioscience Reports</i> , 2021 , 41,	4.1	5
27	Identification and Validation of Key Genes in Hepatocellular Carcinoma by Bioinformatics Analysis. <i>BioMed Research International</i> , 2021 , 2021, 6662114	3	1
26	Sp1-Induced lncRNA Rmrp Promotes Mesangial Cell Proliferation and Fibrosis in Diabetic Nephropathy by Modulating the miR-1a-3p/JunD Pathway. <i>Frontiers in Endocrinology</i> , 2021 , 12, 690784	5.7	1
25	Identification of Key Genes in Gastric Cancer by Bioinformatics Analysis. <i>BioMed Research International</i> , 2020 , 2020, 7658230	3	9
24	Bioinformatics Analysis of Key Genes and circRNA-miRNA-mRNA Regulatory Network in Gastric Cancer. <i>BioMed Research International</i> , 2020 , 2020, 2862701	3	11
23	Construction and analysis of a diabetic nephropathy related protein-protein interaction network reveals nine critical and functionally associated genes. <i>Computational Biology and Chemistry</i> , 2019 , 83, 107115	3.6	1
22	Whole transcriptome analysis of diabetic nephropathy in the db/db mouse model of type 2 diabetes. <i>Journal of Cellular Biochemistry</i> , 2019 , 120, 17520-17533	4.7	8
21	The topological key lncRNA H2k2 from the ceRNA network promotes mesangial cell proliferation in diabetic nephropathy the miR-449a/b/Trim11/Mek signaling pathway. <i>FASEB Journal</i> , 2019 , 33, 11492-1	1506	16
20	Long non-coding RNA Rpph1 promotes inflammation and proliferation of mesangial cells in diabetic nephropathy via an interaction with Gal-3. <i>Cell Death and Disease</i> , 2019 , 10, 526	9.8	44
19	Identification of key genes in non-small cell lung cancer by bioinformatics analysis. <i>PeerJ</i> , 2019 , 7, e821	53.1	12
18	Salvianolic acid B inhibits Ang II-induced VSMC proliferation in vitro and intimal hyperplasia in vivo by downregulating miR-146a expression. <i>Phytomedicine</i> , 2019 , 58, 152754	6.5	11

LIST OF PUBLICATIONS

17	The circRNA circAGFG1 acts as a sponge of miR-195-5p to promote triple-negative breast cancer progression through regulating CCNE1 expression. <i>Molecular Cancer</i> , 2019 , 18, 4	42.1	198
16	LincRNA 1700020I14Rik alleviates cell proliferation and fibrosis in diabetic nephropathy via miR-34a-5p/Sirt1/HIF-1&ignaling. <i>Cell Death and Disease</i> , 2018 , 9, 461	9.8	78
15	Primed atypical ductal hyperplasia-associated fibroblasts promote cell growth and polarity changes of transformed epithelium-like breast cancer MCF-7 cells via miR-200b/c-IKKI ignaling. <i>Cell Death and Disease</i> , 2018 , 9, 122	9.8	10
14	Similar bowtie structures and distinct largest strong components are identified in the transcriptional regulatory networks of Arabidopsis thaliana during photomorphogenesis and heat shock. <i>BioSystems</i> , 2018 , 168, 1-7	1.9	2
13	The Long Noncoding RNA 150Rik Promotes Mesangial Cell Proliferation via miR-451/IGF1R/p38 MAPK Signaling in Diabetic Nephropathy. <i>Cellular Physiology and Biochemistry</i> , 2018 , 51, 1410-1428	3.9	18
12	LincRNA-Gm4419 knockdown ameliorates NF- B /NLRP3 inflammasome-mediated inflammation in diabetic nephropathy. <i>Cell Death and Disease</i> , 2017 , 8, e2583	9.8	151
11	iNOS-derived peroxynitrite mediates high glucose-induced inflammatory gene expression in vascular smooth muscle cells through promoting KLF5 expression and nitration. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2017 , 1863, 2821-2834	6.9	15
10	A Novel Regulatory Mechanism of Smooth Muscle EActin Expression by NRG-1/circACTA2/miR-548f-5p Axis. <i>Circulation Research</i> , 2017 , 121, 628-635	15.7	75
9	miR-451 suppresses the NF-kappaB-mediated proinflammatory molecules expression through inhibiting LMP7 in diabetic nephropathy. <i>Molecular and Cellular Endocrinology</i> , 2016 , 433, 75-86	4.4	73
8	Naringenin Ameliorated Kidney Injury through Let-7a/TGFBR1 Signaling in Diabetic Nephropathy. <i>Journal of Diabetes Research</i> , 2016 , 2016, 8738760	3.9	35
7	Let-7a suppresses cell proliferation via the TGF-ISMAD signaling pathway in cervical cancer. <i>Oncology Reports</i> , 2016 , 36, 3275-3282	3.5	23
6	Promoter hypermethylation of let-7a-3 is relevant to its down-expression in diabetic nephropathy by targeting UHRF1. <i>Gene</i> , 2015 , 570, 57-63	3.8	36
5	Effect of light intensity on survival, growth, and photosynthetic pigment of young seedlings of eelgrass Zostera marina Linnaeus, 1753 (Alismatales: Zosteraceae). <i>Marine Biology Research</i> , 2014 , 10, 745-754	1	5
4	Endonasal endoscopic treatment of recurrent dacryocystitis. <i>Cell Biochemistry and Biophysics</i> , 2013 , 67, 1441-4	3.2	5
3	c-Ski activates cancer-associated fibroblasts to regulate breast cancer cell invasion. <i>Molecular Oncology</i> , 2013 , 7, 1116-28	7.9	34
2	Biological characteristics and genetic heterogeneity between carcinoma-associated fibroblasts and their paired normal fibroblasts in human breast cancer. <i>PLoS ONE</i> , 2013 , 8, e60321	3.7	59
1	MiRNA expression analysis of cancer-associated fibroblasts and normal fibroblasts in breast cancer. <i>International Journal of Biochemistry and Cell Biology</i> , 2012 , 44, 2051-9	5.6	96