

# Qiwei Yang

## List of Publications by Year in descending order

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39  
papers

893  
citations

516710

16  
h-index

501196

28  
g-index

41  
all docs

41  
docs citations

41  
times ranked

1387  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comprehensive Analysis of the Expression and Clinical Significance of THO Complex Members in Hepatocellular Carcinoma. <i>International Journal of General Medicine</i> , 2022, Volume 15, 2695-2713.	1.8	1
2	miR-98-5p protects against cerebral ischemia/reperfusion injury through anti-apoptosis and anti-oxidative stress in mice. <i>Journal of Biochemistry</i> , 2021, 169, 195-206.	1.7	21
3	Expression of TMEM16A in Colorectal Cancer and Its Correlation With Clinical and Pathological Parameters. <i>Frontiers in Oncology</i> , 2021, 11, 652262.	2.8	3
4	Circular RNAs: Expression, localization, and therapeutic potentials. <i>Molecular Therapy</i> , 2021, 29, 1683-1702.	8.2	72
5	Circular RNA CDR1as promotes adipogenic and suppresses osteogenic differentiation of BMSCs in steroid-induced osteonecrosis of the femoral head. <i>Bone</i> , 2020, 133, 115258.	2.9	78
6	Downregulation of BACH1 Protects AGAINST Cerebral Ischemia/Reperfusion Injury through the Functions of HO-1 and NQO1. <i>Neuroscience</i> , 2020, 436, 154-166.	2.3	19
7	Circular RNA hsa_circRNA_0007334 is Predicted to Promote MMP7 and COL1A1 Expression by Functioning as a miRNA Sponge in Pancreatic Ductal Adenocarcinoma. <i>Journal of Oncology</i> , 2019, 2019, 1-16.	1.3	58
8	Exosomal Circular RNA as a Biomarker Platform for the Early Diagnosis of Immune-Mediated Demyelinating Disease. <i>Frontiers in Genetics</i> , 2019, 10, 860.	2.3	34
9	Identification of Key Genes and Circular RNAs in Human Gastric Cancer. <i>Medical Science Monitor</i> , 2019, 25, 2488-2504.	1.1	40
10	Integrated bioinformatics analysis of miRNA expression in Ewing sarcoma and potential regulatory effects of miR-21 via targeting ALCAM/CD166. <i>Artificial Cells, Nanomedicine and Biotechnology</i> , 2019, 47, 2114-2122.	2.8	12
11	Potential Regulatory Effects of miR-182-3p in Osteosarcoma via Targeting EBF2. <i>BioMed Research International</i> , 2019, 2019, 1-9.	1.9	9
12	Network Analyses of Differentially Expressed Genes in Osteoarthritis to Identify Hub Genes. <i>BioMed Research International</i> , 2019, 2019, 1-9.	1.9	15
13	Current Status of Functional Studies on Circular RNAs in Bladder Cancer and their Potential Role as Diagnostic and Prognostic Biomarkers: A Review. <i>Medical Science Monitor</i> , 2019, 25, 3425-3434.	1.1	15
14	The expression of chondrogenesis-related and arthritis-related genes in human ONFH cartilage with different Ficat stages. <i>PeerJ</i> , 2019, 7, e6306.	2.0	12
15	Fisher linear discriminant analysis for classification and prediction of genomic susceptibility to stomach and colorectal cancers based on six STR loci in a northern Chinese Han population. <i>PeerJ</i> , 2019, 7, e7004.	2.0	3
16	Enrichment and Identification of Fetal Nucleated Red Blood Cells from Maternal Blood with Magnetic Nanoparticles and Quantum Dots. <i>Nanoscience and Nanotechnology Letters</i> , 2019, 11, 38-46.	0.4	3
17	Mutation Status and Immunohistochemical Correlation of KRAS, NRAS, and BRAF in 260 Chinese Colorectal and Gastric Cancers. <i>Frontiers in Oncology</i> , 2018, 8, 487.	2.8	29
18	Targeted gene therapy of the HSV-TK/hIL-12 fusion gene controlled by the hSLPI gene promoter of human non-small cell lung cancer in $\frac{1}{2}$ vitro. <i>Oncology Letters</i> , 2018, 15, 6503-6512.	1.8	6

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19	MicroRNA Expression Profiling of Bone Marrow Mesenchymal Stem Cells in Steroid-Induced Osteonecrosis of the Femoral Head Associated with Osteogenesis. <i>Medical Science Monitor</i> , 2018, 24, 1813-1825.	1.1	26
20	Circular RNAs hsa_circ_0032462, hsa_circ_0028173, hsa_circ_0005909 are predicted to promote CADM1 expression by functioning as miRNAs sponge in human osteosarcoma. <i>PLoS ONE</i> , 2018, 13, e0202896.	2.5	33
21	Integration of Gene Expression Profile Data to Screen and Verify Hub Genes Involved in Osteoarthritis. <i>BioMed Research International</i> , 2018, 2018, 1-10.	1.9	22
22	LncRNA expression profiling of BMSCs in osteonecrosis of the femoral head associated with increased adipogenic and decreased osteogenic differentiation. <i>Scientific Reports</i> , 2018, 8, 9127.	3.3	45
23	Ginkgo Biloba L. Extract Reduces H <sub>2</sub> O <sub>2</sub> -Induced Bone Marrow Mesenchymal Stem Cells Cytotoxicity by Regulating Mitogen-Activated Protein Kinase (MAPK) Signaling Pathways and Oxidative Stress. <i>Medical Science Monitor</i> , 2018, 24, 3159-3167.	1.1	13
24	Size-selective separation and overall-amplification of cell-free fetal DNA fragments using PCR-based enrichment. <i>Scientific Reports</i> , 2017, 7, 40936.	3.3	14
25	Association of gene variants of transcription factors PPAR $\gamma$ <sup>3</sup> , RUNX2, Osterix genes and COL2A1, IGFBP3 genes with the development of osteonecrosis of the femoral head in Chinese population. <i>Bone</i> , 2017, 101, 104-112.	2.9	37
26	Validation of reference genes for the normalization of RT-qPCR expression studies in human tongue carcinoma cell lines and tissue. <i>Oncology Letters</i> , 2017, 13, 3951-3957.	1.8	15
27	Endocan silencing induces programmed cell death in hepatocarcinoma. <i>Oncology Letters</i> , 2017, 14, 5333-5339.	1.8	6
28	SPOCK1 promotes the proliferation, migration and invasion of glioma cells through PI3K/AKT and Wnt/ $\beta$ -catenin signaling pathways. <i>Oncology Reports</i> , 2016, 35, 3566-3576.	2.6	54
29	Evaluation and validation of suitable reference genes for reverse transcription-quantitative polymerase chain reaction studies in cholangiocarcinoma patients and cell lines. <i>Oncology Letters</i> , 2016, 11, 2673-2681.	1.8	6
30	Validation of internal reference genes for relative quantitation studies of gene expression in human laryngeal cancer. <i>PeerJ</i> , 2016, 4, e2763.	2.0	6
31	Identification of suitable reference genes for gene expression studies using quantitative polymerase chain reaction in lung cancer in vitro. <i>Molecular Medicine Reports</i> , 2015, 11, 3767-3773.	2.4	34
32	Selection of suitable reference genes for reverse transcription-quantitative polymerase chain reaction analysis of neuronal cells differentiated from bone mesenchymal stem cells. <i>Molecular Medicine Reports</i> , 2015, 12, 2291-2300.	2.4	16
33	Identification of suitable reference genes for investigating gene expression in human gallbladder carcinoma using reverse transcription quantitative polymerase chain reaction. <i>Molecular Medicine Reports</i> , 2015, 11, 2967-2974.	2.4	12
34	Evaluation of suitable control genes for quantitative polymerase chain reaction analysis of maternal plasma cell-free DNA. <i>Molecular Medicine Reports</i> , 2015, 12, 7728-7734.	2.4	5
35	Identification of optimal reference genes for quantitative PCR studies on human mesenchymal stem cells. <i>Molecular Medicine Reports</i> , 2015, 11, 1304-1311.	2.4	29
36	Evaluation of eight reference genes for quantitative polymerase chain reaction analysis in human T lymphocytes co-cultured with mesenchymal stem cells. <i>Molecular Medicine Reports</i> , 2015, 12, 7721-7727.	2.4	8

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37	Validation of suitable reference genes for quantitative polymerase chain reaction analysis in rabbit bone marrow mesenchymal stem cell differentiation. <i>Molecular Medicine Reports</i> , 2015, 12, 2961-2968.	2.4	12
38	Endocan: A new marker for cancer and a target for cancer therapy. <i>Biomedical Reports</i> , 2015, 3, 279-283.	2.0	60
39	Evaluation and validation of the suitable control genes for quantitative PCR studies in plasma DNA for non-invasive prenatal diagnosis. <i>International Journal of Molecular Medicine</i> , 2014, 34, 1681-1687.	4.0	10