Chundi Gao

List of Publications by Citations

42 papers	509 citations	14 h-index	20 g-index
45 ext. papers	768 ext. citations	4.9 avg, IF	3.97 L-index

#	Paper	IF	Citations
42	MicroRNA expression in cervical cancer: Novel diagnostic and prognostic biomarkers. *Journal of Cellular Biochemistry*, **2018**, 119, 7080-7090	4.7	58
41	The construction and analysis of ceRNA networks in invasive breast cancer: a study based on The Cancer Genome Atlas. *Cancer Management and Research*, **2019**, 11, 1-11	3.6	33
40	Exploration of methylation-driven genes for monitoring and prognosis of patients with lung adenocarcinoma. *Cancer Cell International*, **2018**, 18, 194	6.4	31
39	Tumor Characterization in Breast Cancer Identifies Immune-Relevant Gene Signatures Associated With Prognosis. *Frontiers in Genetics*, **2019**, 10, 1119	4.5	29
38	MicroRNAs associated with lung squamous cell carcinoma: New prognostic biomarkers and therapeutic targets. *Journal of Cellular Biochemistry*, **2019**, 120, 18956-18966	4.7	23
37	circRNA-associated ceRNA network construction reveals the circRNAs involved in the progression and prognosis of breast cancer. *Journal of Cellular Physiology*, **2020**, 235, 3973-3983	7	22
36	Prognostic value of aberrantly expressed methylation gene profiles in lung squamous cell carcinoma: A study based on The Cancer Genome Atlas. *Journal of Cellular Physiology*, **2019**, 234, 6519-6528	7	22
35	7-lncRNA Assessment Model for Monitoring and Prognosis of Breast Cancer Patients: Based on Cox Regression and Co-expression Analysis. *Frontiers in Oncology*, **2019**, 9, 1348	5.3	19
34	A Systems Biology-Based Approach to Uncovering Molecular Mechanisms Underlying Effects of Traditional Chinese Medicine Qingdai in Chronic Myelogenous Leukemia, Involving Integration of Network Pharmacology and Molecular Docking Technology. *Medical Science Monitor*, **2018**, 24, 4305-4316	3.2	19
33	Identifying the Antiproliferative Effect of Polysaccharides on Breast Cancer: Coupling Network Pharmacology With Targetable Screening From the Cancer Genome Atlas. *Frontiers in Oncology*, **2019**, 9, 368	5.3	17
32	The Modulatory Properties of Treatment on Triple-Negative Breast Cancer: An Integrated Pharmacological Method. *Frontiers in Pharmacology*, **2019**, 10, 1171	5.6	17
31	Identification of the key pathways and genes involved in HER2-positive breast cancer with brain metastasis. *Pathology Research and Practice*, **2019**, 215, 152475	3.4	16
30	Four lncRNAs associated with breast cancer prognosis identified by coexpression network analysis. *Journal of Cellular Physiology*, **2019**, 234, 14019-14030	7	16
29	SNP mutation-related genes in breast cancer for monitoring and prognosis of patients: A study based on the TCGA database. *Cancer Medicine*, **2019**, 8, 2303-2312	4.8	14
28	Exploring the Mechanism of Danshen against Myelofibrosis by Network Pharmacology and Molecular Docking. *Evidence-based Complementary and Alternative Medicine*, **2018**, 2018, 8363295	2.3	14
27	Cardiotoxicity of doxorubicin-based cancer treatment: What is the protective cognition that phytochemicals provide us?. *Pharmacological Research*, **2020**, 160, 105062	10.2	13
26	A review of the biological activity and pharmacology of cryptotanshinone, an important active constituent in Danshen. *Biomedicine and Pharmacotherapy*, **2021**, 137, 111332	7.5	13

25	DNA methylation-based diagnostic and prognostic biomarkers of nonsmoking lung adenocarcinoma patients. *Journal of Cellular Biochemistry*, **2019**, 120, 13520-13530	4.7	11
24	Comprehensive analysis of prognostic biomarkers in lung adenocarcinoma based on aberrant lncRNA-miRNA-mRNA networks and Cox regression models. *Bioscience Reports*, **2020**, 40,	4.1	11
23	Target Analysis and Mechanism of Podophyllotoxin in the Treatment of Triple-Negative Breast Cancer. *Frontiers in Pharmacology*, **2020**, 11, 1211	5.6	11
22	Deciphering Key Pharmacological Pathways of Qingdai Acting on Chronic Myeloid Leukemia Using a Network Pharmacology-Based Strategy. *Medical Science Monitor*, **2018**, 24, 5668-5688	3.2	11
21	A four-mRNA model to improve the prediction of breast cancer prognosis. *Gene*, **2019**, 721, 144100	3.8	10
20	Identification of prognostic biomarkers for breast cancer based on miRNA and mRNA co-expression network. *Journal of Cellular Biochemistry*, **2019**, 120, 15378-15388	4.7	9
19	Natural Polysaccharides and Their Derivates: A Promising Natural Adjuvant for Tumor Immunotherapy. *Frontiers in Pharmacology*, **2021**, 12, 621813	5.6	8
18	Construction and Analysis of Competing Endogenous RNA Networks for Breast Cancer Based on TCGA Dataset. *BioMed Research International*, **2020**, 2020, 4078596	3	7
17	Tumor Mutation Burden and Immune Invasion Characteristics in Triple Negative Breast Cancer: Genome High-Throughput Data Analysis. *Frontiers in Immunology*, **2021**, 12, 650491	8.4	7
16	Identification of key candidate targets and pathways for the targeted treatment of leukemia stem cells of chronic myelogenous leukemia using bioinformatics analysis. *Molecular Genetics & Genomic Medicine*, **2019**, 7, e851	2.3	6
15	Biomarker expression analysis in different age groups revealed age was a risk factor for breast cancer. *Journal of Cellular Physiology*, **2020**, 235, 4268-4278	7	6
14	The association between statin use and endometrial cancer survival outcome: A meta-analysis. *Medicine (United States)*, **2018**, 97, e13264	1.8	6
13	Identification of key candidate genes and miRNA-mRNA target pairs in chronic lymphocytic leukemia by integrated bioinformatics analysis. *Molecular Medicine Reports*, **2019**, 19, 362-374	2.9	5
12	Developing DNA methylation-based prognostic biomarkers of acute myeloid leukemia. *Journal of Cellular Biochemistry*, **2018**, 119, 10041-10050	4.7	4
11	A Novel Glycolysis-Related Four-mRNA Signature for Predicting the Survival of Patients With Breast Cancer. *Frontiers in Genetics*, **2021**, 12, 606937	4.5	3
10	Efficacy and safety of de-escalation bone- modifying agents for cancer patients with bone metastases: a systematic review and meta-analysis. *Cancer Management and Research*, **2018**, 10, 3809-3823	3.6	3
9	Deciphering the mechanism of Indirubin and its derivatives in the inhibition of Imatinib resistance using a "drug target prediction-gene microarray analysis-protein network construction" strategy. *BMC Complementary and Alternative Medicine*, **2019**, 19, 75	4.7	2
8	6-lncRNA Assessment Model for Monitoring and Prognosis of HER2-Positive Breast Cancer: Based on Transcriptome Data. *Pathology and Oncology Research*, **2021**, 27, 609083	2.6	2

7	An mRNA characterization model predicting survival in patients with invasive breast cancer based on The Cancer Genome Atlas database. *Cancer Biomarkers*, **2021**, 30, 417-428	3.8	2
6	Development of a risk scoring system for evaluating the prognosis of patients with Her2-positive breast cancer. *Cancer Cell International*, **2020**, 20, 121	6.4	2
5	Cryptotanshinone Is a Intervention for ER-Positive Breast Cancer: An Integrated Approach to the Study of Natural Product Intervention Mechanisms. *Frontiers in Pharmacology*, **2020**, 11, 592109	5.6	2
4	Determination of Genetic and Epigenetic Modifications-Related Prognostic Biomarkers of Breast Cancer: Genome High-Throughput Data Analysis. *Journal of Oncology*, **2021**, 2021, 2143362	4.5	2
3	Gene signatures of 6-methyladenine regulators in women with lung adenocarcinoma and development of a risk scoring system: a retrospective study using the cancer genome atlas database. *Aging*, **2021**, 13, 3957-3968	5.6	1
2	Natural compounds: A new perspective on targeting polarization and infiltration of tumor-associated macrophages in lung cancer.. *Biomedicine and Pharmacotherapy*, **2022**, 151, 113096	7.5	1
1	Development of a Novel Immune-Related Gene Prognostic Index for Breast Cancer.. *Frontiers in Immunology*, **2022**, 13, 845093	8.4	1