

Cun Liu

List of Publications by Year in descending order

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Version: 2024-02-01

56
papers

1,160
citations

361045

20
h-index

454577

30
g-index

59
all docs

59
docs citations

59
times ranked

1557
citing authors

#	ARTICLE	IF	CITATIONS
1	MicroRNA expression in cervical cancer: Novel diagnostic and prognostic biomarkers. <i>Journal of Cellular Biochemistry</i> , 2018, 119, 7080-7090.	1.2	82
2	Tumor Characterization in Breast Cancer Identifies Immune-Relevant Gene Signatures Associated With Prognosis. <i>Frontiers in Genetics</i> , 2019, 10, 1119.	1.1	64
3	Exploration of methylation-driven genes for monitoring and prognosis of patients with lung adenocarcinoma. <i>Cancer Cell International</i> , 2018, 18, 194.	1.8	59
4	The construction and analysis of ceRNA networks in invasive breast cancer: a study based on The Cancer Genome Atlas. <i>Cancer Management and Research</i> , 2019, Volume 11, 1-11.	0.9	49
5	A review of the biological activity and pharmacology of cryptotanshinone, an important active constituent in Danshen. <i>Biomedicine and Pharmacotherapy</i> , 2021, 137, 111332.	2.5	47
6	Meta-analysis of the association between the dietary inflammatory index (DII) and breast cancer risk. <i>European Journal of Clinical Nutrition</i> , 2019, 73, 509-517.	1.3	46
7	7-lncRNA Assessment Model for Monitoring and Prognosis of Breast Cancer Patients: Based on Cox Regression and Co-expression Analysis. <i>Frontiers in Oncology</i> , 2019, 9, 1348.	1.3	34
8	MicroRNAs associated with lung squamous cell carcinoma: New prognostic biomarkers and therapeutic targets. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 18956-18966.	1.2	33
9	The Modulatory Properties of Astragalus membranaceus Treatment on Triple-Negative Breast Cancer: An Integrated Pharmacological Method. <i>Frontiers in Pharmacology</i> , 2019, 10, 1171.	1.6	32
10	Impact of endometriosis on risk of ovarian, endometrial and cervical cancers: a meta-analysis. <i>Archives of Gynecology and Obstetrics</i> , 2019, 299, 35-46.	0.8	32
11	SNP mutation-related genes in breast cancer for monitoring and prognosis of patients: A study based on the TCGA database. <i>Cancer Medicine</i> , 2019, 8, 2303-2312.	1.3	31
12	Prognostic value of aberrantly expressed methylation gene profiles in lung squamous cell carcinoma: A study based on The Cancer Genome Atlas. <i>Journal of Cellular Physiology</i> , 2019, 234, 6519-6528.	2.0	31
13	Integrative transcriptome data mining for identification of core lncRNAs in breast cancer. <i>PeerJ</i> , 2019, 7, e7821.	0.9	30
14	Tanshinone IIA: New Perspective on the Anti-Tumor Mechanism of A Traditional Natural Medicine. <i>The American Journal of Chinese Medicine</i> , 2022, 50, 209-239.	1.5	30
15	Four lncRNAs associated with breast cancer prognosis identified by coexpression network analysis. <i>Journal of Cellular Physiology</i> , 2019, 234, 14019-14030.	2.0	29
16	circRNA-associated ceRNA network construction reveals the circRNAs involved in the progression and prognosis of breast cancer. <i>Journal of Cellular Physiology</i> , 2020, 235, 3973-3983.	2.0	29
17	Tumor Mutation Burden and Immune Invasion Characteristics in Triple Negative Breast Cancer: Genome High-Throughput Data Analysis. <i>Frontiers in Immunology</i> , 2021, 12, 650491.	2.2	29
18	Role of Flavonoids in the Treatment of Iron Overload. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 685364.	1.8	29

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19	Identifying the Antiproliferative Effect of Astragalus Polysaccharides on Breast Cancer: Coupling Network Pharmacology With Targetable Screening From the Cancer Genome Atlas. <i>Frontiers in Oncology</i> , 2019, 9, 368.	1.3	27
20	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. <i>Medical Science Monitor</i> , 2018, 24, 4305-4316.	0.5	25
21	Identification of the key pathways and genes involved in HER2-positive breast cancer with brain metastasis. <i>Pathology Research and Practice</i> , 2019, 215, 152475.	1.0	24
22	Cardiotoxicity of doxorubicin-based cancer treatment: What is the protective cognition that phytochemicals provide us?. <i>Pharmacological Research</i> , 2020, 160, 105062.	3.1	24
23	Natural Polysaccharides and Their Derivates: A Promising Natural Adjuvant for Tumor Immunotherapy. <i>Frontiers in Pharmacology</i> , 2021, 12, 621813.	1.6	22
24	Nanoformulations of Ursolic Acid: A Modern Natural Anticancer Molecule. <i>Frontiers in Pharmacology</i> , 2021, 12, 706121.	1.6	22
25	DNA methylation-based diagnostic and prognostic biomarkers of nonsmoking lung adenocarcinoma patients. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 13520-13530.	1.2	21
26	Exploring the Mechanism of Danshen against Myelofibrosis by Network Pharmacology and Molecular Docking. <i>Evidence-based Complementary and Alternative Medicine</i> , 2018, 2018, 1-11.	0.5	18
27	Deciphering Key Pharmacological Pathways of Qingdai Acting on Chronic Myeloid Leukemia Using a Network Pharmacology-Based Strategy. <i>Medical Science Monitor</i> , 2018, 24, 5668-5688.	0.5	17
28	Target Analysis and Mechanism of Podophyllotoxin in the Treatment of Triple-Negative Breast Cancer. <i>Frontiers in Pharmacology</i> , 2020, 11, 1211.	1.6	17
29	Identification of lncRNAs associated with lung squamous cell carcinoma prognosis in the competitive endogenous RNA network. <i>PeerJ</i> , 2019, 7, e7727.	0.9	17
30	Identification of key candidate genes and miRNA-mRNA target pairs in chronic lymphocytic leukemia by integrated bioinformatics analysis. <i>Molecular Medicine Reports</i> , 2019, 19, 362-374.	1.1	16
31	The effect of long noncoding RNAs HOX transcript antisense intergenic RNA single-nucleotide polymorphisms on breast cancer, cervical cancer, and ovarian cancer susceptibility: A meta-analysis. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 7056-7067.	1.2	16
32	Biomarker expression analysis in different age groups revealed age was a risk factor for breast cancer. <i>Journal of Cellular Physiology</i> , 2020, 235, 4268-4278.	2.0	14
33	Development of a risk scoring system for evaluating the prognosis of patients with Her2-positive breast cancer. <i>Cancer Cell International</i> , 2020, 20, 121.	1.8	14
34	Remodeling the Epigenetic Landscape of Cancer—Application Potential of Flavonoids in the Prevention and Treatment of Cancer. <i>Frontiers in Oncology</i> , 2021, 11, 705903.	1.3	14
35	Research progress of phenolic compounds regulating IL-6 to exert antitumor effects. <i>Phytotherapy Research</i> , 2021, 35, 6720-6734.	2.8	13
36	The association between statin use and endometrial cancer survival outcome. <i>Medicine (United Tj ETQq0 0 0 rgBT/O</i>	0.4	12

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37	Natural compounds: A new perspective on targeting polarization and infiltration of tumor-associated macrophages in lung cancer. <i>Biomedicine and Pharmacotherapy</i> , 2022, 151, 113096.	2.5	11
38	Comparative efficacy of different targeted therapies plus fulvestrant for advanced breast cancer following progression on prior endocrine therapy: a network meta-analysis. <i>Cancer Management and Research</i> , 2018, Volume 10, 5869-5880.	0.9	10
39	Identification of key candidate targets and pathways for the targeted treatment of leukemia stem cells of chronic myelogenous leukemia using bioinformatics analysis. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e851.	0.6	10
40	A Novel Glycolysis-Related Four-mRNA Signature for Predicting the Survival of Patients With Breast Cancer. <i>Frontiers in Genetics</i> , 2021, 12, 606937.	1.1	10
41	Construction and Analysis of Competing Endogenous RNA Networks for Breast Cancer Based on TCGA Dataset. <i>BioMed Research International</i> , 2020, 2020, 1-10.	0.9	9
42	<p>Efficacy and safety of targeted therapy for metastatic HER2-positive breast cancer in the first-line treatment: a Bayesian network meta-analysis<p>. <i>OncoTargets and Therapy</i> , 2019, Volume 12, 959-974.	1.0	8
43	Proteinâ€protein interaction networks and different clustering analysis in Burkittâ€™s lymphoma. <i>Hematology</i> , 2018, 23, 391-398.	0.7	6
44	Efficacy and safety of de-escalation bone-modifying agents for cancer patients with bone metastases: a systematic review and meta-analysis. <i>Cancer Management and Research</i> , 2018, Volume 10, 3809-3823.	0.9	6
45	6-lncRNA Assessment Model for Monitoring and Prognosis of HER2-Positive Breast Cancer: Based on Transcriptome Data. <i>Pathology and Oncology Research</i> , 2021, 27, 609083.	0.9	6
46	An mRNA characterization model predicting survival in patients with invasive breast cancer based on The Cancer Genome Atlas database. <i>Cancer Biomarkers</i> , 2021, 30, 417-428.	0.8	5
47	Developing DNA methylationâ€based prognostic biomarkers of acute myeloid leukemia. <i>Journal of Cellular Biochemistry</i> , 2018, 119, 10041-10050.	1.2	4
48	Deciphering the mechanism of Indirubin and its derivatives in the inhibition of Imatinib resistance using a â€drug target prediction-gene microarray analysis-protein network constructionâ€strategy. <i>BMC Complementary and Alternative Medicine</i> , 2019, 19, 75.	3.7	4
49	Deciphering of Key Pharmacological Pathways of Poria Cocos Intervention in Breast Cancer Based on Integrated Pharmacological Method. <i>Evidence-based Complementary and Alternative Medicine</i> , 2020, 2020, 1-11.	0.5	4
50	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. <i>Aging</i> , 2021, 13, 3957-3968.	1.4	4
51	Identifying the Effect of Ursolic Acid Against Triple-Negative Breast Cancer: Coupling Network Pharmacology With Experiments Verification. <i>Frontiers in Pharmacology</i> , 2021, 12, 685773.	1.6	4
52	Acupuncture for Aromatase Inhibitorâ€Related Joint Pain Among Breast Cancer Patients. <i>JAMA - Journal of the American Medical Association</i> , 2018, 320, 2270.	3.8	3
53	Cryptotanshinone Is a Intervention for ER-Positive Breast Cancer: An Integrated Approach to the Study of Natural Product Intervention Mechanisms. <i>Frontiers in Pharmacology</i> , 2020, 11, 592109.	1.6	3
54	Determination of Genetic and Epigenetic Modifications-Related Prognostic Biomarkers of Breast Cancer: Genome High-Throughput Data Analysis. <i>Journal of Oncology</i> , 2021, 2021, 1-12.	0.6	3

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55	Analysis of the expression patterns and clinical relevance of m6A regulators in 33 cancer types. <i>Future Oncology</i> , 2022, 18, 565-577.	1.1	1
56	Letter to the editor: efficacy and safety of a combination of HER2-targeted agents as first-line treatment for metastatic HER2-positive breast cancer: a network meta-analysis. <i>Expert Opinion on Drug Safety</i> , 2018, 17, 1151-1152.	1.0	0