

# Chundi Gao

## List of Publications by Year in Descending Order

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**Version:** 2024-04-10

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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.

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

| #  | Paper  | IF   | Citations |
|----|--|------|-----------|
| 42 | Natural compounds: A new perspective on targeting polarization and infiltration of tumor-associated macrophages in lung cancer.. <i>Biomedicine and Pharmacotherapy</i> , <b>2022</b> , 151, 113096                                | 7.5  | 1         |
| 41 | Development of a Novel Immune-Related Gene Prognostic Index for Breast Cancer.. <i>Frontiers in Immunology</i> , <b>2022</b> , 13, 845093  | 8.4  | 1         |
| 40 | Tumor Mutation Burden and Immune Invasion Characteristics in Triple Negative Breast Cancer: Genome High-Throughput Data Analysis. <i>Frontiers in Immunology</i> , <b>2021</b> , 12, 650491  | 8.4  | 7         |
| 39 | 6-lncRNA Assessment Model for Monitoring and Prognosis of HER2-Positive Breast Cancer: Based on Transcriptome Data. <i>Pathology and Oncology Research</i> , <b>2021</b> , 27, 609083  | 2.6  | 2         |
| 38 | An mRNA characterization model predicting survival in patients with invasive breast cancer based on The Cancer Genome Atlas database. <i>Cancer Biomarkers</i> , <b>2021</b> , 30, 417-428   | 3.8  | 2         |
| 37 | Natural Polysaccharides and Their Derivates: A Promising Natural Adjuvant for Tumor Immunotherapy. <i>Frontiers in Pharmacology</i> , <b>2021</b> , 12, 621813   | 5.6  | 8         |
| 36 | A review of the biological activity and pharmacology of cryptotanshinone, an important active constituent in Danshen. <i>Biomedicine and Pharmacotherapy</i> , <b>2021</b> , 137, 111332   | 7.5  | 13        |
| 35 | 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> , <b>2021</b> , 13, 3957-3968 | 5.6  | 1         |
| 34 | A Novel Glycolysis-Related Four-mRNA Signature for Predicting the Survival of Patients With Breast Cancer. <i>Frontiers in Genetics</i> , <b>2021</b> , 12, 606937   | 4.5  | 3         |
| 33 | Determination of Genetic and Epigenetic Modifications-Related Prognostic Biomarkers of Breast Cancer: Genome High-Throughput Data Analysis. <i>Journal of Oncology</i> , <b>2021</b> , 2021, 2143362                               | 4.5  | 2         |
| 32 | Comprehensive analysis of prognostic biomarkers in lung adenocarcinoma based on aberrant lncRNA-miRNA-mRNA networks and Cox regression models. <i>Bioscience Reports</i> , <b>2020</b> , 40,                                       | 4.1  | 11        |
| 31 | circRNA-associated ceRNA network construction reveals the circRNAs involved in the progression and prognosis of breast cancer. <i>Journal of Cellular Physiology</i> , <b>2020</b> , 235, 3973-3983                                | 7    | 22        |
| 30 | Biomarker expression analysis in different age groups revealed age was a risk factor for breast cancer. <i>Journal of Cellular Physiology</i> , <b>2020</b> , 235, 4268-4278   | 7    | 6         |
| 29 | Cardiotoxicity of doxorubicin-based cancer treatment: What is the protective cognition that phytochemicals provide us?. <i>Pharmacological Research</i> , <b>2020</b> , 160, 105062  | 10.2 | 13        |
| 28 | Construction and Analysis of Competing Endogenous RNA Networks for Breast Cancer Based on TCGA Dataset. <i>BioMed Research International</i> , <b>2020</b> , 2020, 4078596   | 3    | 7         |
| 27 | Target Analysis and Mechanism of Podophyllotoxin in the Treatment of Triple-Negative Breast Cancer. <i>Frontiers in Pharmacology</i> , <b>2020</b> , 11, 1211  | 5.6  | 11        |
| 26 | Development of a risk scoring system for evaluating the prognosis of patients with Her2-positive breast cancer. <i>Cancer Cell International</i> , <b>2020</b> , 20, 121   | 6.4  | 2         |

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| 25 | 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> , <b>2020</b> , 11, 592109   | 5.6  | 2  |
| 24 | A four-mRNA model to improve the prediction of breast cancer prognosis. <i>Gene</i> , <b>2019</b> , 721, 144100   | 3.8  | 10 |
| 23 | MicroRNAs associated with lung squamous cell carcinoma: New prognostic biomarkers and therapeutic targets. <i>Journal of Cellular Biochemistry</i> , <b>2019</b> , 120, 18956-18966   | 4.7  | 23 |
| 22 | Identification of the key pathways and genes involved in HER2-positive breast cancer with brain metastasis. <i>Pathology Research and Practice</i> , <b>2019</b> , 215, 152475  | 3.4  | 16 |
| 21 | Identifying the Antiproliferative Effect of Polysaccharides on Breast Cancer: Coupling Network Pharmacology With Targetable Screening From the Cancer Genome Atlas. <i>Frontiers in Oncology</i> , <b>2019</b> , 9, 368   | 5.3  | 17 |
| 20 | Identification of prognostic biomarkers for breast cancer based on miRNA and mRNA co-expression network. <i>Journal of Cellular Biochemistry</i> , <b>2019</b> , 120, 15378-15388   | 4.7  | 9  |
| 19 | 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. <i>BMC Complementary and Alternative Medicine</i> , <b>2019</b> , 19, 75 | 4.7  | 2  |
| 18 | DNA methylation-based diagnostic and prognostic biomarkers of nonsmoking lung adenocarcinoma patients. <i>Journal of Cellular Biochemistry</i> , <b>2019</b> , 120, 13520-13530   | 4.7  | 11 |
| 17 | SNP mutation-related genes in breast cancer for monitoring and prognosis of patients: A study based on the TCGA database. <i>Cancer Medicine</i> , <b>2019</b> , 8, 2303-2312   | 4.8  | 14 |
| 16 | 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 &amp; Genomic Medicine</i> , <b>2019</b> , 7, e851                                | 2.3  | 6  |
| 15 | The Modulatory Properties of Treatment on Triple-Negative Breast Cancer: An Integrated Pharmacological Method. <i>Frontiers in Pharmacology</i> , <b>2019</b> , 10, 1171  | 5.6  | 17 |
| 14 | 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> , <b>2019</b> , 9, 1348  | 5.3  | 19 |
| 13 | Tumor Characterization in Breast Cancer Identifies Immune-Relevant Gene Signatures Associated With Prognosis. <i>Frontiers in Genetics</i> , <b>2019</b> , 10, 1119   | 4.5  | 29 |
| 12 | Four lncRNAs associated with breast cancer prognosis identified by coexpression network analysis. <i>Journal of Cellular Physiology</i> , <b>2019</b> , 234, 14019-14030  | 7    | 16 |
| 11 | 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> , <b>2019</b> , 234, 6519-6528  | 7.28 | 22 |
| 10 | 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> , <b>2019</b> , 11, 1-11   | 3.6  | 33 |
| 9  | Identification of key candidate genes and miRNA-mRNA target pairs in chronic lymphocytic leukemia by integrated bioinformatics analysis. <i>Molecular Medicine Reports</i> , <b>2019</b> , 19, 362-374  | 2.9  | 5  |
| 8  | MicroRNA expression in cervical cancer: Novel diagnostic and prognostic biomarkers. <i>Journal of Cellular Biochemistry</i> , <b>2018</b> , 119, 7080-7090  | 4.7  | 58 |

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|---|--|-----|----|
| 7 | The association between statin use and endometrial cancer survival outcome: A meta-analysis. <i>Medicine (United States)</i> , <b>2018</b> , 97, e13264  | 1.8 | 6  |
| 6 | Deciphering Key Pharmacological Pathways of Qingdai Acting on Chronic Myeloid Leukemia Using a Network Pharmacology-Based Strategy. <i>Medical Science Monitor</i> , <b>2018</b> , 24, 5668-5688   | 3.2 | 11 |
| 5 | 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> , <b>2018</b> , 24, 4305-4316 | 3.2 | 19 |
| 4 | Exploring the Mechanism of Danshen against Myelofibrosis by Network Pharmacology and Molecular Docking. <i>Evidence-based Complementary and Alternative Medicine</i> , <b>2018</b> , 2018, 8363295   | 2.3 | 14 |
| 3 | Exploration of methylation-driven genes for monitoring and prognosis of patients with lung adenocarcinoma. <i>Cancer Cell International</i> , <b>2018</b> , 18, 194  | 6.4 | 31 |
| 2 | 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> , <b>2018</b> , 10, 3809-3823   | 3.6 | 3  |
| 1 | Developing DNA methylation-based prognostic biomarkers of acute myeloid leukemia. <i>Journal of Cellular Biochemistry</i> , <b>2018</b> , 119, 10041-10050   | 4.7 | 4  |