

# Shan Lu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7064336/publications.pdf>

Version: 2024-02-01

17  
papers

115  
citations

1684188

5  
h-index

1474206

9  
g-index

19  
all docs

19  
docs citations

19  
times ranked

77  
citing authors

#	ARTICLE	IF	CITATIONS
1	A novel strategy to reveal clinical advantages and molecular mechanism of aidi injection in the treatment of pancreatic cancer based on network meta-analysis and network pharmacology. <i>Journal of Ethnopharmacology</i> , 2022, 285, 114852.	4.1	5
2	Exploring the Multicomponent Synergy Mechanism of Yinzhihuang Granule in Inhibiting Inflammation-Cancer Transformation of Hepar Based on Integrated Bioinformatics and Network Pharmacology. <i>BioMed Research International</i> , 2022, 2022, 1-13.	1.9	1
3	An advanced network pharmacology study to explore the novel molecular mechanism of Compound Kushen Injection for treating hepatocellular carcinoma by bioinformatics and experimental verification. <i>BMC Complementary Medicine and Therapies</i> , 2022, 22, 54.	2.7	3
4	Revealing the Mechanism of Huazhi Rougan Granule in the Treatment of Nonalcoholic Fatty Liver Through Intestinal Flora Based on 16S rRNA, Metagenomic Sequencing and Network Pharmacology. <i>Frontiers in Pharmacology</i> , 2022, 13, 875700.	3.5	4
5	Investigation on the Efficiency of Chinese Herbal Injections combined with Concurrent Chemoradiotherapy for Treating Nasopharyngeal Carcinoma based on Multidimensional Bayesian Network Meta-analysis. <i>Frontiers in Pharmacology</i> , 2021, 12, 656724.	3.5	4
6	Dissecting the novel mechanism of reduning injection in treating Coronavirus Disease 2019 (COVID-19) based on network pharmacology and experimental verification. <i>Journal of Ethnopharmacology</i> , 2021, 273, 113871.	4.1	23
7	Investigation on the mechanisms of guiqi huoxue capsule for treating cervical spondylosis based on network pharmacology and molecular docking. <i>Medicine (United States)</i> , 2021, 100, e26643.	1.0	4
8	An Advanced Systems Pharmacology Strategy Reveals AKR1B1, MMP2, PTGER3 as Key Genes in the Competing Endogenous RNA Network of Compound Kushen Injection Treating Gastric Carcinoma by Integrated Bioinformatics and Experimental Verification. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 742421.	3.7	4
9	Identification of Key Genes Associated With the Process of Hepatitis B Inflammation and Cancer Transformation by Integrated Bioinformatics Analysis. <i>Frontiers in Genetics</i> , 2021, 12, 654517.	2.3	8
10	Biodata Mining of Differentially Expressed Genes between Acute Myocardial Infarction and Unstable Angina Based on Integrated Bioinformatics. <i>BioMed Research International</i> , 2021, 2021, 1-19.	1.9	5
11	A Multidimensional Bayesian Network Meta-Analysis of Chinese Herbal Injections for Treating Non-small Cell Lung Cancer With Gemcitabine and Cisplatin. <i>Frontiers in Pharmacology</i> , 2021, 12, 739673.	3.5	4
12	High Throughput Transcriptome Data Analysis and Computational Verification Reveal Immunotherapy Biomarkers of Compound Kushen Injection for Treating Triple-Negative Breast Cancer. <i>Frontiers in Oncology</i> , 2021, 11, 747300.	2.8	2
13	Investigation on the clinical efficacy and mechanism of compound kushen injection in treating esophageal cancer based on multi-dimensional network meta-analysis and in vitro experiment. <i>Journal of Ethnopharmacology</i> , 2021, 279, 114386.	4.1	8
14	A network pharmacology approach to reveal the pharmacological targets and biological mechanism of compound kushen injection for treating pancreatic cancer based on WGCNA and in vitro experiment validation. <i>Chinese Medicine</i> , 2021, 16, 121.	4.0	21
15	A New Strategy to Identify ceRNA-Based CCDC144NL-AS1/SERPINE1 Regulatory Axis as a Novel Prognostic Biomarker for Stomach Adenocarcinoma via High Throughput Transcriptome Data Mining and Computational Verification. <i>Frontiers in Oncology</i> , 2021, 11, 802727.	2.8	4
16	Exploring potential mechanisms of Suhexiang Pill against COVID-19 based on network pharmacology and molecular docking. <i>Medicine (United States)</i> , 2021, 100, e27112.	1.0	6
17	Identification of candidate biomarkers correlated with the pathogenesis and prognosis of breast cancer via integrated bioinformatics analysis. <i>Medicine (United States)</i> , 2020, 99, e23153.	1.0	9