

# Luo Mei

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2482452/publications.pdf>

Version: 2024-02-01

11  
papers

730  
citations

1163117  
8  
h-index

1281871  
11  
g-index

11  
all docs

11  
docs citations

11  
times ranked

830  
citing authors

#	ARTICLE	IF	CITATIONS
1	ImmuCellAI: A Unique Method for Comprehensive T Cell Subsets Abundance Prediction and its Application in Cancer Immunotherapy. <i>Advanced Science</i> , 2020, 7, 1902880.	11.2	558
2	ImmuCellAI-mouse: a tool for comprehensive prediction of mouse immune cell abundance and immune microenvironment depiction. <i>Bioinformatics</i> , 2022, 38, 785-791.	4.1	53
3	Leukemia cell-derived microvesicles induce T cell exhaustion via miRNA delivery. <i>Oncotarget</i> , 2018, 7, e1448330.	4.6	24
4	Differential Co-expression and Regulatory Network Analysis Uncover the Relapse Factor and Mechanism of T Cell Acute Leukemia. <i>Molecular Therapy - Nucleic Acids</i> , 2018, 12, 184-194.	5.1	20
5	FFLtool: a web server for transcription factor and miRNA feed forward loop analysis in human. <i>Bioinformatics</i> , 2020, 36, 2605-2607.	4.1	19
6	Integrating Transcriptome and Experiments Reveals the Anti-diabetic Mechanism of Cyclocarya paliurus Formula. <i>Molecular Therapy - Nucleic Acids</i> , 2018, 13, 419-430.	5.1	12
7	Systematic Transcriptome and Regulatory Network Analyses Reveal the Hypoglycemic Mechanism of <i>Dendrobium fimbriatum</i> . <i>Molecular Therapy - Nucleic Acids</i> , 2020, 19, 1-14.	5.1	11
8	Genetic, Pharmacogenomic, and Immune Landscapes of Enhancer RNAs Across Human Cancers. <i>Cancer Research</i> , 2022, 82, 785-790.	0.9	11
9	Investigating the Molecular Mechanism of Aqueous Extract of <i>Cyclocarya paliurus</i> on Ameliorating Diabetes by Transcriptome Profiling. <i>Frontiers in Pharmacology</i> , 2018, 9, 912.	3.5	9
10	Regulatory network analysis reveals the oncogenesis roles of feed-forward loops and therapeutic target in T-cell acute lymphoblastic leukemia. <i>BMC Medical Genomics</i> , 2019, 12, 8.	1.5	7
11	CCLA: an accurate method and web server for cancer cell line authentication using gene expression profiles. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	6