

# Ya-Ru Miao

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3344816/publications.pdf>

Version: 2024-02-01

14  
papers

1,957  
citations

840776

11  
h-index

1058476

14  
g-index

14  
all docs

14  
docs citations

14  
times ranked

2842  
citing authors

#	ARTICLE	IF	CITATIONS
1	EVAtlas: a comprehensive database for ncRNA expression in human extracellular vesicles. <i>Nucleic Acids Research</i> , 2022, 50, D111-D117.	14.5	27
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	A comprehensive platelet expression atlas (PEA) resource and platelet transcriptome landscape. <i>American Journal of Hematology</i> , 2022, 97, E18.	4.1	3
4	Case Report: Multi-Omics Analysis and CAR-T Treatment of a Chronic Myeloid Leukemia Blast Crisis Case 5 Years After the Discontinuation of TKI. <i>Frontiers in Oncology</i> , 2021, 11, 739871.	2.8	7
5	A 6-Membrane Protein Gene score for prognostic prediction of cytogenetically normal acute myeloid leukemia in multiple cohorts. <i>Journal of Cancer</i> , 2020, 11, 251-259.	2.5	12
6	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
7	hTFtarget: A Comprehensive Database for Regulations of Human Transcription Factors and Their Targets. <i>Genomics, Proteomics and Bioinformatics</i> , 2020, 18, 120-128.	6.9	201
8	SNP2APA: a database for evaluating effects of genetic variants on alternative polyadenylation in human cancers. <i>Nucleic Acids Research</i> , 2020, 48, D226-D232.	14.5	37
9	ImmuCellAI: A Unique Method for Comprehensive Tâ€Cell Subsets Abundance Prediction and its Application in Cancer Immunotherapy. <i>Advanced Science</i> , 2020, 7, 1902880.	11.2	558
10	Identification of STAB1 in Multiple Datasets as a Prognostic Factor for Cytogenetically Normal AML: Mechanism and Drug Indications. <i>Molecular Therapy - Nucleic Acids</i> , 2019, 18, 476-484.	5.1	22
11	AnimalTFDB 3.0: a comprehensive resource for annotation and prediction of animal transcription factors. <i>Nucleic Acids Research</i> , 2019, 47, D33-D38.	14.5	592
12	EVmiRNA: a database of miRNA profiling in extracellular vesicles. <i>Nucleic Acids Research</i> , 2019, 47, D89-D93.	14.5	209
13	SEGreg: a database for human specifically expressed genes and their regulations in cancer and normal tissue. <i>Briefings in Bioinformatics</i> , 2019, 20, 1322-1328.	6.5	18
14	lncRNASNP2: an updated database of functional SNPs and mutations in human and mouse lncRNAs. <i>Nucleic Acids Research</i> , 2018, 46, D276-D280.	14.5	199