



## (Core Project R03OD038391)

### Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD038391-01	Leveraging Heterogenous Common Fund Data Sets and Beyond for Identifying Lung Cancer Subtypes	\$307K	9 publications	0 repositories	0 properties

## Publications

Published works associated with this project.

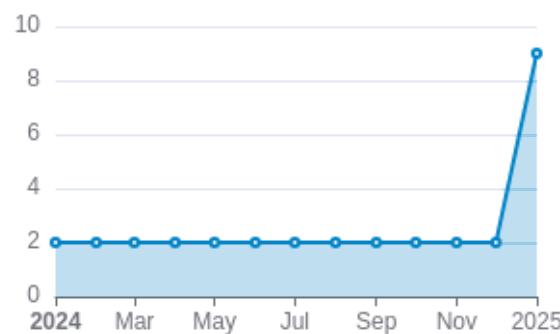
ID	Title	Author s	RC R	SJ R	Cita tion s	Cit. /ye ar	Journal	Pub lish ed	Upd ated
<a href="#">39573886</a>	SAMP: Identifying antimicrobial peptides by an ensemble learning model based on proportionalized ...	Feng, Junxi ...6 more... Wan, Shibiao	0. 46 5	0. 8 6	2	2	Briefings in functional genomics	2024	Dec 1, 2025
<a href="#">40313658</a>	A Comprehensive Review on RNA Subcellular Localization Prediction.	Zhang, Cece ...3 more... Wan, Shibiao	0	0	0	0	ArXiv	2025	Dec 1, 2025
<a href="#">40060540</a>	Accurate identification of medulloblastoma subtypes from diverse data sources with severe batch e...	Sun, Mengta o ...1 more... Wan, Shibiao	0	0	0	0	bioRxiv : the preprint server for biology	2025	Dec 1, 2025

<a href="#">41246237</a>	RanBALL: An Ensemble Machine Learning Framework for Accurate Subtype Identification of Pediatric ...	Li, Lushen g ...5 more... Wan, Shibiao	1. 0 1 0 0 4	Advanced intelligent systems (Weinheim an der Bergstrasse, Germany)	2025	Dec 1, 2025
<a href="#">39436320</a>	A review of artificial intelligence-based brain age estimation and its applications for related d...	Azzam, Moha med ...6 more... Wang, Jieqion g	0. 0 8 3 3 6	Briefings in functional genomics	2025	Dec 1, 2025
<a href="#">40475602</a>	AttentionAML: An Attention-based Deep Learning Framework for Accurate Molecular Categorization of...	Li, Lushen g ...2 more... Wan, Shibiao	0 0 0 0 0	bioRxiv : the preprint server for biology	2025	Dec 1, 2025
<a href="#">39386448</a>	RanBALL: An Ensemble Random Projection Model for Identifying Subtypes of B-Cell Acute Lymphoblast...	Li, Lushen g ...5 more...	0 0 2 2	bioRxiv : the preprint server for biology	2025	Dec 1, 2025

		Wan, Shibiao					
<a href="#">39386613</a>	WIMOAD: Weighted Integration of Multi-Omics Data with Meta Learning for Alzheimer's Disease Diagn...	Xiao, Hanyu ...1 more...	0	0	1	1	bioRxiv : the preprint server for biology
		Wan, Shibiao					
<a href="#">39605468</a>	Functional Connectivity Alterations in Cocaine Use Disorder: Insights from the Triple Network Mod...	Xu, Ziyang ...4 more...	0	0	0	0	bioRxiv : the preprint server for biology
		Wang, Jieqiong					

### Publications (cumulative)

Total: 9



## Notes

RCR [Relative Citation Ratio ↗](#)

SJR [Scimago Journal Rank ↗](#)

## </> Repositories

Software repositories associated with this project.

Name	Description	Last Commit	S	F	W	C	Issues	PRs	I	P	R	C	o	C	n	o	L	C	L
			t	o	a	m			A	v	d	t	e	r	i	c	o	a	n
			g	s	r	k	s	s	m	u	b	f	n	e	r	i	o	o	u

No data

## Notes

Repository For storing, tracking changes to, and collaborating on a piece of software.

PR "Pull request", a draft change (new feature, bug fix, etc.) to a repo.

Closed/Open Resolved/unresolved.

Issue/PR Avg Average time issues/pull requests stay open for before being closed.

Only the main/default branch is considered for metrics like # of commits.

# of dependencies is totaled from all manifests in repo, direct and transitive, e.g. package.json + package-lock.json .



Website metrics associated with this project.

## Notes

Active Users [Distinct users who visited the website ↗](#).

New Users [Users who visited the website for the first time ↗](#).

Engaged Sessions [Visits that had significant interaction ↗](#).

"Top" metrics are measured by number of engaged sessions.