



## (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	15 publications	0 repositories	0 properties

## Publications

Published works associated with this project.

ID	Title	Author s	RC R	SJ R	Cita tions	Cit. /ye ar	Journal	Pub lished	Upda ted
<a href="#">39271266</a> 	Multi-omics based artificial intelligence for cancer research.	Li, Lushen g ...2 more... Wan, Shibiao	3. 77 9	0	16	16	Adv Cancer Res	2024	Dec 28, 2025
<a href="#">39573886</a> 	SAMP: Identifying antimicrobial peptides by an ensemble learning model based on proportionalized ...	Feng, Junxi ...6 more... Wan, Shibiao	0. 43 1	0. 8 6	2	2	Briefings in functional genomics	2024	Dec 28, 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	0. 8 6	3	3	Briefings in functional genomics	2025	Dec 28, 2025

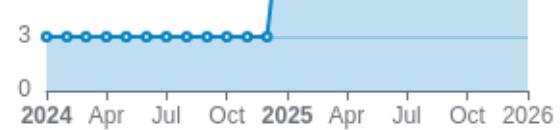
<a href="#">39386613</a>	WIMOAD: Weighted Integration of Multi-Omics Data with Meta Learning for Alzheimer's Disease Diagn...	Xiao, Hanyu ...1 more... Wan, Shibiao	0    0    2    2	bioRxiv : the preprint server for biology	2025	Dec 28, 2025
<a href="#">39386448</a>	RanBALL: An Ensemble Random Projection Model for Identifying Subtypes of B-Cell Acute Lymphoblast...	Li, Lushen g ...5 more... Wan, Shibiao	0    0    2    2	bioRxiv : the preprint server for biology	2025	Dec 28, 2025
<a href="#">39605468</a>	Functional Connectivity Alterations in Cocaine Use Disorder: Insights from the Triple Network Mod...	Xu, Ziyang ...4 more... Wang, Jieqiong	0    0    0    0	bioRxiv : the preprint server for biology	2024	Dec 28, 2025
<a href="#">41509387</a>	MOTLAB: A Weighted Multi-Omics Transfer Learning Approach to Mitigate Breast Cancer Racial Dispar...	Baek, Min- Jeong ...3 more... Wan, Shibiao	0    0    0    0	bioRxiv : the preprint server for biology	2025	Jan 23, 2026

<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	bioRxiv : the preprint server for biology	2025	Dec 28, 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 28, 2025
<a href="#">41347120</a>	RPSLearner: A Novel Approach Based on Random Projection and Deep Stacking Learning for Categorizi...	Wu, Xincha o ...1 more... Wan, Shibiao	1.0    0    1    0    4	Advanced intelligent systems (Weinheim an der Bergstrasse, Germany)	2025	Dec 28, 2025

### Publications (cumulative)

Total: 15





## Notes

RCR [Relative Citation Ratio ↗](#)

SJR [Scimago Journal Rank ↗](#)

# </> Repositories

Software repositories associated with this project

Built on Jan 26, 2026

Developed with support from NIH Award [U54 OD036472](#)

Name	Description	Last Commit	S	F	W	C	Issues	PRs	I	P	R	E	n	O	L	C	L	
			t	o	a	o			s	R	a	i	d	r	e	c	o	a
bioRxiv	bioRxiv is a preprint server for life sciences research. It is a community-driven platform that allows researchers to share their work before it has been peer-reviewed. The repository contains the source code for the bioRxiv website and various tools used in the preprint submission process.	2023-01-26T14:45:23Z	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10

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.

## Analytics

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.