



(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	12 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
39271266 	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
39573886 	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
39386613 	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

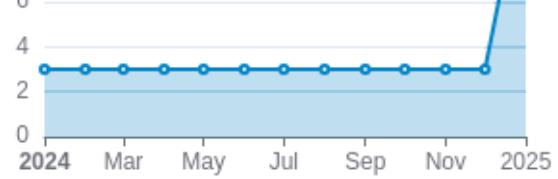
39436320	A review of artificial intelligence-based brain age estimation and its applications for related d...	Azzam, Mohamed ...6 more... Wang, Jieqiong	0. 8 3 3	Briefings in functional genomics	2025	Dec 28, 2025		
40313658	A Comprehensive Review on RNA Subcellular Localization Prediction.	Zhang, Cece ...3 more... Wan, Shibiao	0 0 0 0	ArXiv	2025	Dec 28, 2025		
41246237	RanBALL: An Ensemble Machine Learning Framework for Accurate Subtype Identification of Pediatric ...	Li, Lushen g ...5 more... Wan, Shibiao	1. 1 0 0	Advanced intelligent systems (Weinheim an der Bergstrasse, Germany)	2025	Jan 21, 2026		
40060540	Accurate identification of medulloblastoma subtypes from diverse data sources with severe batch e...	Sun, Mengtao ...1 more... Wan, Shibiao	0 0 0 0	bioRxiv : the preprint server for biology	2025	Dec 28, 2025		

39605468	Functional Connectivity Alterations in Cocaine Use Disorder: Insights from the Triple Network Mod...	Xu, Ziyang ...4 more... Wang, Jieqion g	0 0 0 0	bioRxiv : the preprint server for biology	2024	Dec 28, 2025
39386448	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
40475602	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

Publications (cumulative)

Total: 12





Notes

RCR [Relative Citation Ratio](#) ↗

SJR [Scimago Journal Rank](#) ↗

</> Repositories

Software repositories associated with this project

Built on Jan 23, 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 provides a platform for researchers to share their work before it has been peer-reviewed and published in a journal. The repository contains the source code for the bioRxiv website and various tools used in the preprint submission process.	2023-12-01T12:00:00Z	1000	1000	1000	1000	1000	1000	1000	1000	1000	1000	1000	1000	1000	1000	1000	1000

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.