



(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
39436320 	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

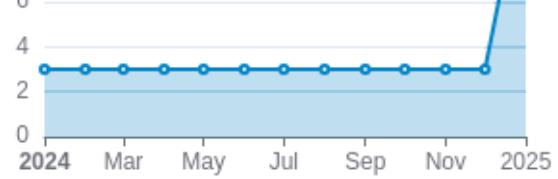
40313658	A Comprehensive Review on RNA Subcellular Localization Prediction.	Zhang, Cece ...3 more... Wan, Shibiao	0 0 0 0	ArXiv	2025	Dec 28, 2025
39386613	WIMOAD: Weighted Integration of Multi-Omics Data with Meta Learning for Alzheimer's Disease Diagn...	Xiao, Hanyu ...1 more... Wan, Shibiao	0 0 2 2	bioRxiv : the preprint server for biology	2025	Dec 28, 2025
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
41446223	A Multi-Modal Transfer Learning Framework to Reduce Health Disparities in Prostate Adenocarcinoma.	Li, Lushen g ...1 more... Wan, Shibiao	0 0 0 0	bioRxiv : the preprint server for biology	2025	Jan 6, 2026

41246237	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 28, 2025
40060540	Accurate identification of medulloblastoma subtypes from diverse data sources with severe batch e...	Sun, Mengta o ...1 more... Wan, Shibiao	0 0 0 0 0	bioRxiv : the preprint server for biology	2025	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

Publications (cumulative)

Total: 12





Notes

RCR [Relative Citation Ratio](#) ↗

SJR [Scimago Journal Rank](#) ↗

</> Repositories

Software repositories associated with this project

Built on Jan 13, 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.	2023-12-01T12:00:00Z	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.