



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

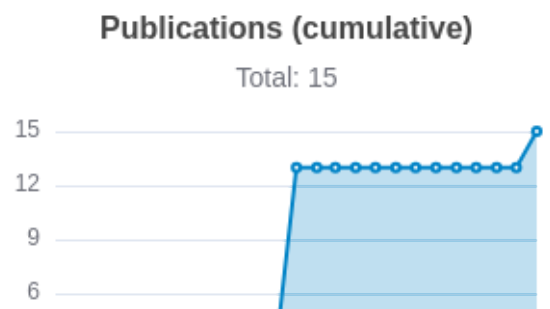
 Publications

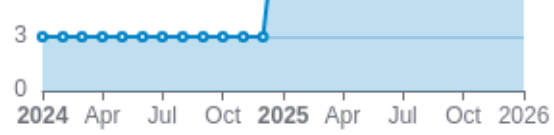
Published works associated with this project.

ID	Title	Author s	RC R	SJ R	Cita tion s	Cit. /ye ar	Journal	Pub lish ed	Upda ted
39271266  DOI 	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  DOI 	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  DOI 	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 6	3	3	Briefings in functional genomics	2025	Dec 28, 2025

39386613  DOI 	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
39386448  DOI 	RanBALL: An Ensemble Random Projection Model for Identifying Subtypes of B-Cell Acute Lymphoblast...	Li, Lusheng ...5 more... Wan, Shibiao	0	0	2	2	bioRxiv : the preprint server for biology	2025	Dec 28, 2025
39605468  DOI 	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
41509387  DOI 	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

40475602  DOI 	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
40313658  DOI 	A Comprehensive Review on RNA Subcellular Localization Prediction.	Zhang, Cece ...3 more... Wan, Shibiao	0	0	0	0	ArXiv	2025	Dec 28, 2025
41347120  DOI 	RPSLearner: A Novel Approach Based on Random Projection and Deep Stacking Learning for Categorizi...	Wu, Xinchao o...1 more... Wan, Shibiao	0	14	0	0	Advanced intelligent systems (Weinheim an der Bergstrasse, Germany)	2025	Dec 28, 2025





Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

</> Repositories

Software repositories associated with this project

Built on Jan 31, 2026

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

Name	Description	Tags	Last Commit	Stars	Forks	Watchers	Commits	Issues	PRs	Issue Avg	PR Avg	Readme	Contributing	Code of Con.	License	Contribs	Languages

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.