



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.

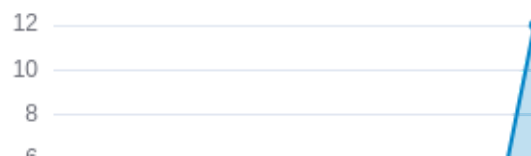
ID	Title	Author s	RC R	SJ R	Cita tion s	Cit. /ye ar	Journal	Pub lish ed	Upda ted
39271266  DOI 	Multi-omics based artificial intelligence for cancer research.	Li, Lushen g ...2 more... Wan, Shibiao	3. 77 9	0	16	16	Adv Cancer Res	2024	Dec 28, 2025
39573886  DOI 	SAMP: Identifying antimicrobial peptides 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
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

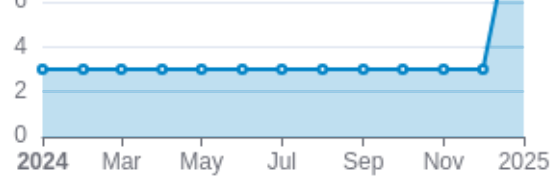
39436320  DOI 	A review of artificial intelligence-based brain age estimation and its applications for related d...	Azzam, Mohamed ...6 more... Wang, Jieqiong	0	0.86	3	3	Briefings in functional genomics	2025	Dec 28, 2025
40313658  DOI 	A Comprehensive Review on RNA Subcellular Localization Prediction.	Zhang, Cece ...3 more... Wan, Shibiao	0	0	0	0	ArXiv	2025	Dec 28, 2025
41246237  DOI 	RanBALL: An Ensemble Machine Learning Framework for Accurate Subtype Identification of Pediatric ...	Li, Lusheng ...5 more... Wan, Shibiao	0	1.14	0	0	Advanced intelligent systems (Weinheim an der Bergstrasse, Germany)	2025	Jan 21, 2026
40060540  DOI 	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  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
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
40475602  DOI 	AttentionAML: An Attention-based Deep Learning Framework for Accurate Molecular Categorization of...	Li, Lusheng ...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 21, 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.