



# 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	9 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	Upd ated
<a href="#">39573886</a>  <a href="#">DOI</a> 	SAMP: Identifying antimicrobial peptides by an ensemble learning model based on proportionalized ...	Feng, Junxi ...6 more... Wan, Shibiao	0. 46 5	0. 8 6	2	2	Briefings in functional genomics	2024	Dec 1, 2025
<a href="#">40313658</a>  <a href="#">DOI</a> 	A Comprehensive Review on RNA Subcellular Localization Prediction.	Zhang, Cece ...3 more... Wan, Shibiao	0	0	0	0	ArXiv	2025	Dec 1, 2025
<a href="#">40060540</a>  <a href="#">DOI</a> 	Accurate identification of medulloblastoma subtypes from diverse data sources with severe batch e...	Sun, Mengta o ...1 more... Wan, Shibiao	0	0	0	0	bioRxiv : the preprint server for biology	2025	Dec 1, 2025

<a href="#">41246237</a>  <a href="#">DOI</a> 	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	Dec 1, 2025
<a href="#">39436320</a>  <a href="#">DOI</a> 	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 1, 2025
<a href="#">40475602</a>  <a href="#">DOI</a> 	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 1, 2025
<a href="#">39386448</a>  <a href="#">DOI</a> 	RanBALL: An Ensemble Random Projection Model for Identifying Subtypes of B-Cell Acute Lymphoblast...	Li, Lusheng ...5 more...	0	0	2	2	bioRxiv : the preprint server for biology	2025	Dec 1, 2025

		Wan, Shibiao									
<a href="#">39386613</a> 	<a href="#">DOI</a> 	WIMOAD: Weighted Integration of Multi-Omics Data with Meta Learning for Alzheimer's Disease Diagn...	Xiao, Hanyu ...1 more... Wan, Shibiao	0	0	1	1	bioRxiv : the preprint server for biology	2025	Dec 1, 2025	
<a href="#">39605468</a> 	<a href="#">DOI</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 1, 2025	



Notes

RCR [Relative Citation Ratio](#)  
SJR [Scimago Journal Rank](#)

</> Repositories

Software repositories associated with this project.

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



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.