



## (Core Project R03OD038391)

### Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories
1R03OD038391-01	Leveraging Heterogenous Common Fund Data Sets and Beyond for Identifying Lung Cancer Subtypes	\$307,000.00	9 publications	0 repositories

### Analytics

0 properties

## Publications

Published works associated with this project.

ID	Title	Authors	RC R	SJ R	Cita tions	Cit./ yea r	Journal	Publ ishe d	Updat ed
<a href="#">39573886</a>	SAMP: Identifying antimicrobial peptides by an ensemble learning model based on proportionalized ...	Feng, Junxi ...6 more... Wan, Shibiao	0.4 65	0	2	2	Brief Funct Genomics	2024	Dec 10, 2025 (just now)
<a href="#">40060540</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	2025	Dec 10, 2025 (just now)
<a href="#">40313658</a>	A Comprehensive Review on RNA Subcellular Localization Prediction.	Zhang, Cece ...3 more... Wan, Shibiao	0	0	0	0	ArXiv	2025	Dec 10, 2025 (just now)

<a href="#">41246237</a>	RanBALL: An Ensemble Machine Learning Framework for Accurate Subtype Identification of Pediatric ...	Li, Lusheng ...5 more... Wan, Shibiao	0    0    0    0	Adv Intell Syst	2025	Dec 10, 2025 (just now)
<a href="#">39436320</a>	A review of artificial intelligence-based brain age estimation and its applications for related d...	Azzam, Moham ed ...6 more... Wang, Jieqiong	0    0    3    3	Brief Funct Genomics	2025	Dec 10, 2025 (just now)
<a href="#">40475602</a>	AttentionAML: An Attention-based Deep Learning Framework for Accurate Molecular Categorization of...	Li, Lusheng ...2 more... Wan, Shibiao	0    0    0    0	bioRxiv	2025	Dec 10, 2025 (just now)
<a href="#">39386448</a>	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	2025	Dec 10, 2025 (just now)

[39386613](#) 

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 2025 2025 (just now)

[39605468](#) 

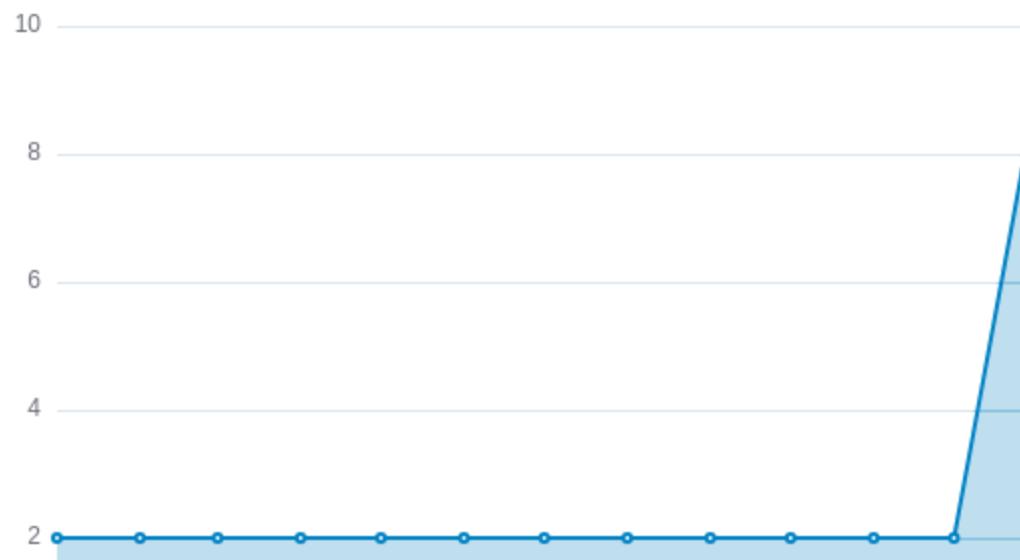
Functional Connectivity Alterations in Cocaine Use Disorder: Insights from the Triple Network Mod...

Xu,  
Ziyang  
...4  
more...  
Wang,  
Jieqiong

0 0 0 0 bioRxiv 2024 2025 (just now)

### Publications (cumulative)

Total: 9





Software repositories associated with this project.

## Notes

RCR [Relative Citation Ratio](#) ↗

Name	Description	Tags	Last Commit	Stars	Forks	Watchers	Commits	Issues	PRs
No data									

Name	Issue Avg	PR Avg	Readme	Contributing	Code of Con.	License	Contrib.	Languages	Dependencies
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.

Built on Dec 10, 2025

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

## 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.