



(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 |
| 40313658  | A Comprehensive Review on RNA Subcellular Localization Prediction. | Zhang, Cece ...3 more... Wan, Shibiao | 0 | 0 | 0 | 0 | ArXiv | 2025 | Dec 28, 2025 |

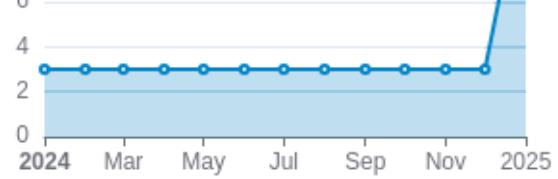
| | | | | | | |
|--------------------------|--|---|-----------------------------|---|------|--------------|
| 39436320 | A review of artificial intelligence-based brain age estimation and its applications for related d... | Azzam, Moha med ...6 more... Wang, Jieqion g | 0. 0 8 3 3 6 | Briefings in functional genomics | 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, Mengtao ...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 17, 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 |

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.