

L Core Project R030D038391

O Details

| Projects | Name | Award | Publications | Repositories | Analytics |
|-----------------|-----------------------------------------------------------------------------------------------------|--------------|----------------|----------------|--------------|
| 1R03OD038391-01 | Leveraging Heterogenous Common Fund Data Sets and Beyond for Identifying Lung Cancer Subtypes | \$307,000.00 | 8 publications | 0 repositories | 0 properties |

Publications

Published works associated with this project.

| ID | Title | Authors | R C R | SJ R | Cita tion s | Cit./ yea r | Journal | Publ ishe d | Updat ed |
|------------|------------------------------------------------------------------------------------|----------------|-------------|---------|-------------------|-------------------|---------|-------------------|-------------|
| 40060540 🗹 | Accurate identification of medulloblastoma subtypes from diverse data sources with | Sun, Mengta | 0 | 0 | 0 | 0 | bioRxiv | 2025 | Aug 26, |

| | and the state of t | | | | | | | | 2025 |
|-----------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------|---|---|---|---|----------------------------|------|-------------------------------------|
| | severe batch e | o 1 more Wan, Shibiao | | | | | | | 2025 (just now) |
| 39436320 2 DOI 2 | A review of artificial intelligence-based brain age estimation and its applications for related d | Azzam, Moham ed 6 more Wang, Jieqiong | 0 | 0 | 2 | 2 | Brief Funct Genomics | 2025 | Aug 26, 2025 (just now) |
| 39386448 乙 DOI 乙 | 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 | Aug 26, 2025 (just now) |
| 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 | 0 | 0 | bioRxiv | 2025 | Aug 26, 2025 (just now) |
| 39573886 🗹 | SAMP: Identifying antimicrobial peptides by an ensemble learning model based on | Feng, Junxi | 0 | 0 | 2 | 2 | Brief Funct | 2024 | Aug 26, |

| | proportionalized | 6 more Wan, Shibiao | | | | | Genomics | | 2025 (just now) |
|---------------------|---------------------------------------------------------------------------------------------------------|------------------------------------------------|---|---|---|---|----------|------|-------------------------------------|
| 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 | 2025 | Aug 26, 2025 (just now) |
| 39605468 🗹 DOI 🗹 | Functional Connectivity Alterations in Cocaine Use Disorder: Insights from the Triple Network Mod | Xu, Ziyang4 more Wang, Jieqiong | 0 | 0 | 0 | 0 | bioRxiv | 2024 | Aug 26, 2025 (just now) |
| 40313658 ☑ DOI ☑ | A Comprehensive Review on RNA Subcellular Localization Prediction. | Zhang, Cece 3 more Wan, Shibiao | 0 | 0 | 0 | 0 | ArXiv | 2025 | Aug 26, 2025 (just now) |

Notes

Publications (cumulative)

Total: 8



| Name | Description | Stars | Watchers | Forks | Issues | PRs | Commits | Contrib. | |
|---------|-------------|-------|----------|-------|--------|-----|---------|----------|--|
| No data | | | | | | | | | |

| Name | Tags | Last Commit | Avg Issue | Avg PR | Languages | License | Readme | Contributing | Dependencies |
|------|------|-------------|-----------|--------|-----------|---------|--------|--------------|--------------|
| | | | | | No data | | | | |

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.

Avg Issue/PR 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

Traffic metrics of websites associated with this project.

Notes

Active Users <u>Distinct users who visited the website</u> 2.

New Users <u>Users who visited the website for the first time</u> **?**.

Engaged Sessions <u>Visits that had significant interaction</u> 2.

"Top" metrics are measured by number of engaged sessions.