



## (Core Project R03OD034493)

### Overview

High-level info about this project.

| Projects        | Name  | Award  | Publications   | Repositories   | Analytics    |
|-----------------|---|--------|----------------|----------------|--------------|
| 1R03OD034493-01 | Cross Repository Metabolomics Data and Workflow Integration | \$303K | 8 publications | 0 repositories | 0 properties |

## Publications

Published works associated with this project.

| ID   | Title   | Authors  | RC<br>R   | SJR       | Cita<br>tion<br>s | Cit.<br>/ye<br>ar | Journal          | Pub<br>lished | Upd<br>ated       |
|--|---|--|-----------|-----------|-------------------|-------------------|------------------|---------------|-------------------|
| <a href="#">38471500</a>    | The underappreciated diversity of bile acid modifications.                          | Mohanty,<br>Ipsita<br>...26 more...<br>Dorrestein,<br>Pieter C | 27.<br>13 | 22.<br>61 | 130               | 65                | Cell             | 202<br>4      | Feb<br>1,<br>2026 |
| <a href="#">38769143</a>    | Reproducible mass spectrometry data processing and compound annotation in MZmine 3. | Heuckeroth,<br>Steffen<br>...17 more...<br>Pluskal,<br>Tomáš   | 19.<br>49 | 5.8<br>54 | 62                | 31                | Nature protocols | 202<br>4      | Feb<br>1,<br>2026 |
| <a href="#">40355727</a>  | A universal language for finding mass spectrometry data patterns.                   | Damiani,<br>Tito<br>...72 more...<br>Wang,<br>Mingxun          | 13.<br>40 | 17.<br>25 | 27                | 27                | Nature methods   | 202<br>5      | Feb<br>1,<br>2026 |
| <a href="#">40021805</a>  | A guide to reverse metabolomics-a framework for big data discovery strategy.        | Charron-<br>Lamoureux,<br>Vincent<br>...13 more...             | 5.4<br>69 | 5.8<br>54 | 11                | 11                | Nature protocols | 202<br>5      | Feb<br>1,<br>2026 |

|                          |  |  |           |                |    |           |   |  |          |                   |  |
|--------------------------|--|--|-----------|----------------|----|-----------|---|--|----------|-------------------|--|
|                          | Dorresteijn,<br>Pieter C   |  |           |                |    |           |   |  |          |                   |  |
| <a href="#">40020666</a> | The International Space Station has a unique and extreme microbial and chemical environment drive... | Salido,<br>Rodolfo A<br>...23 more...              | 5.0<br>11 | 22.<br>61<br>2 | 10 | 10        | Cell  |  | 202<br>5 | Feb<br>1,<br>2026 |  |
| <a href="#">38918178</a> | Leptochelins A-C, Cytotoxic Metallophores Produced by Geographically Dispersed <i>Leptothoe</i> S... | Avalon,<br>Nicole E<br>...28 more...               | 4.9<br>17 | 5.5<br>54      | 17 | 8.5       | Journal of the American Chemical Society              |  | 202<br>4 | Feb<br>1,<br>2026 |  |
| <a href="#">38830143</a> | ModiFinder: Tandem Mass Spectral Alignment Enables Structural Modification Site Localization.        | Shahneh,<br>Mohammad<br>Reza Zare<br>...10 more... | 3.5<br>54 | 0.7<br>04      | 15 | 7.5       | Journal of the American Society for Mass Spectrometry |  | 202<br>4 | Feb<br>1,<br>2026 |  |
| <a href="#">36688502</a> | Unified and Standardized Mass Spectrometry Data Processing in Python Using spectrum_utils.           | Bittremieux,<br>Wout<br>...5 more...               | 2.2<br>31 | 1.1<br>39      | 16 | 5.3<br>33 | Journal of proteome research                          |  | 202<br>3 | Feb<br>1,<br>2026 |  |

### Publications (cumulative)

Total: 8



## Notes

RCR [Relative Citation Ratio ↗](#)

SJR [Scimago Journal Rank ↗](#)

## </> Repositories

Software repositories associated with this project.

C  
o C  
n o . C L

Built on Feb 2, 2026

Developed with support from NIH Award U54 OD036472

| m<br>e | Description | g<br>s | Last Commit | a<br>r<br>s | r<br>k<br>s | h<br>e<br>s | m<br>it<br>rs | ... | Issues | PRs | ~<br>A<br>v<br>g | A<br>v<br>g | ~<br>d<br>m<br>e | ~<br>b<br>u<br>t | ~<br>f<br>C<br>o | ~<br>e<br>n<br>s | ~<br>r<br>i<br>c | ~<br>a<br>g<br>e |
|--------|-------------|--------|-------------|-------------|-------------|-------------|---------------|-----|--------|-----|------------------|-------------|------------------|------------------|------------------|------------------|------------------|------------------|
|--------|-------------|--------|-------------|-------------|-------------|-------------|---------------|-----|--------|-----|------------------|-------------|------------------|------------------|------------------|------------------|------------------|------------------|

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.

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.