



# Core Project R03OD034493



## Details











| Projects        | Name  | Award        | Publications   | Repositories   | Analytics    |
|-----------------|---|--------------|----------------|----------------|--------------|
| 1R03OD034493-01 | Cross Repository Metabolomics Data and Workflow Integration | \$302,764.00 | 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 | Publ<br>ishe<br>d | Upd<br>ated       |
|---|--|----------------------------------|----------------|----------------|-------------------|-------------------|---------|-------------------|-------------------|
| <a href="#">38471500</a><br><a href="#">DOI</a> | The underappreciated diversity of bile acid modifications. | Mohanty, Ipsita<br>...26 more... | 26.<br>86<br>7 | 22.<br>61<br>2 | 110               | 110               | Cell    | 202<br>4          | Dec<br>3,<br>2025 |

|   |  |   |                |   |    |    |                  |          |                                    |
|---|--|---|----------------|---|----|----|------------------|----------|------------------------------------|
|   |  | Dorrestein,<br>Pieter C   |                |   |    |    |                  |          | (just<br>now)                      |
| <a href="#">38769143</a> <br><a href="#">DOI</a>      | Reproducible mass spectrometry data processing and compound annotation in MZmine 3.                  | Heuckeroth,<br>Steffen<br>...17 more...<br>Pluskal,<br>Tomáš                  | 18.<br>63<br>1 | 0 | 52 | 52 | Nat<br>Protoc    | 202<br>4 | Dec<br>3,<br>2025<br>(just<br>now) |
| <a href="#">40355727</a> <br><a href="#">DOI</a>      | A universal language for finding mass spectrometry data patterns.                                    | Damiani, Tito<br>...72 more...<br>Wang,<br>Mingxun                            | 9.5<br>18      | 0 | 19 | 19 | Nat<br>Methods   | 202<br>5 | Dec<br>3,<br>2025<br>(just<br>now) |
| <a href="#">38918178</a> <br><a href="#">DOI</a>    | Leptochelins A-C, Cytotoxic Metallophores Produced by Geographically Dispersed <i>Leptothoe</i> S... | Avalon,<br>Nicole E<br>...28 more...<br>Gerwick,<br>William H                 | 5.6<br>87      | 0 | 16 | 16 | J Am<br>Chem Soc | 202<br>4 | Dec<br>3,<br>2025<br>(just<br>now) |
| <a href="#">40021805</a> <br><a href="#">DOI</a>  | A guide to reverse metabolomics-a framework for big data discovery strategy.                         | Charron-<br>Lamoureux,<br>Vincent<br>...13 more...<br>Dorrestein,<br>Pieter C | 2.9<br>93      | 0 | 6  | 6  | Nat<br>Protoc    | 202<br>5 | Dec<br>3,<br>2025<br>(just<br>now) |
| <a href="#">38830143</a> <br><a href="#">DOI</a>  | ModiFinder: Tandem Mass Spectral Alignment Enables Structural  | Shahneh,<br>Mohammad  | 2.6<br>53      | 0 | 10 | 10 | J Am Soc<br>Mass | 202<br>4 | Dec<br>3,                          |

|   |  |   |           |                |    |     |                      |          |  |                                    |
|---|--|---|-----------|----------------|----|-----|----------------------|----------|--|------------------------------------|
|   | Modification Site Localization.  | Reza Zare<br>...10 more...<br>Wang,<br>Mingxun                  |           |                |    |     |                      | Spectrom |  | 2025<br>(just<br>now)              |
| <a href="#">36688502</a> <br><a href="#">DOI</a>  | Unified and Standardized Mass Spectrometry Data Processing in Python Using spectrum_utils.           | Bittremieux,<br>Wout<br>...5 more...<br>Dorrestein,<br>Pieter C | 1.9<br>17 | 0              | 13 | 6.5 | J<br>Proteome<br>Res | 2023     |  | Dec<br>3,<br>2025<br>(just<br>now) |
| <a href="#">40020666</a> <br><a href="#">DOI</a>  | The International Space Station has a unique and extreme microbial and chemical environment drive... | Salido,<br>Rodolfo A<br>...23 more...<br>Knight, Rob            | 0         | 22.<br>61<br>2 | 4  | 4   | Cell                 | 2025     |  | Dec<br>3,<br>2025<br>(just<br>now) |

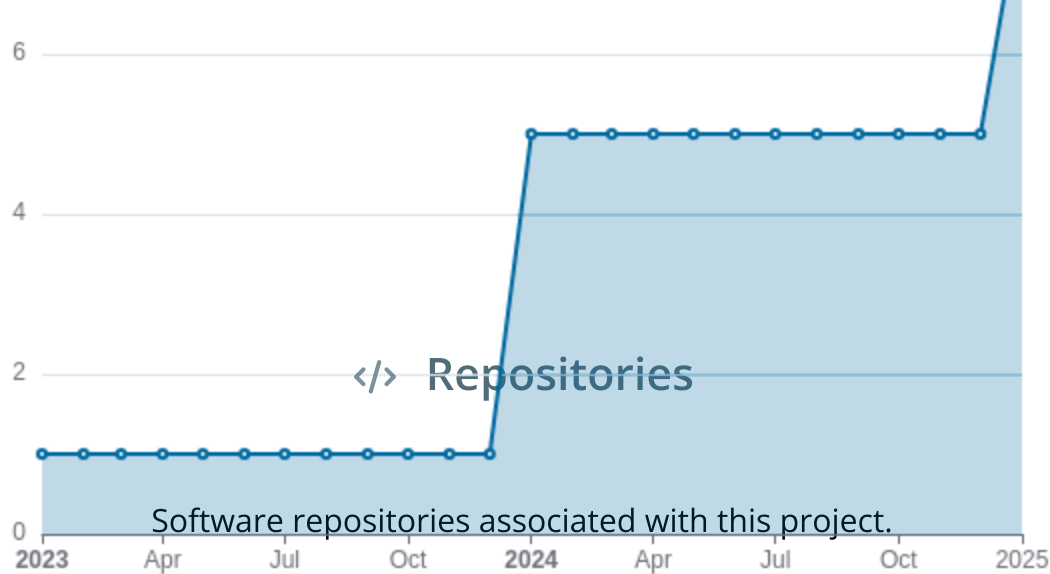
## Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

**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 |      |             |           |        |           |         |        |              |              |

Notes

- RepositoryFor 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/OpenResolved/unresolved.
- Avg Issue/PRAverage 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      [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.

Built on Dec 3, 2025

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