



# Core Project R03OD034501









## Overview







High-level info about this project.

| Projects        | Name   | Award  | Publications   | Repositories   | Analytics    |
|-----------------|--|--------|----------------|----------------|--------------|
| 1R03OD034501-01 | Integration of GTEx and HuBMAP data to gain population-level cell-type-specific insights | \$315K | 6 publications | 0 repositories | 0 properties |

## Publications

Published works associated with this project.

| ID  | Title  | Author<br>s  | RC<br>R       | SJ<br>R       | Cita<br>tion<br>s | Cit.<br>/ye<br>ar | Journal   | Publ<br>ishe<br>d | Upd<br>ated       |
|---|--|--|---------------|---------------|-------------------|-------------------|---|-------------------|-------------------|
| <a href="#">38168620</a> <br><a href="#">DOI</a>      | scMD facilitates cell type deconvolution using single-cell DNA methylation references.               | Cai,<br>Manqi<br>...2<br>more...<br>Wang,<br>Jiebiao             | 9.<br>78<br>7 | 2.<br>07<br>1 | 42                | 42                | Communications biology  | 202<br>4          | Dec<br>1,<br>2025 |
| <a href="#">37563770</a> <br><a href="#">DOI</a>  | Transcriptional risk scores in Alzheimer's disease: From pathology to cognition.                     | Pyun,<br>Jung-<br>Min<br>...7<br>more...<br>Nho,<br>Kwang<br>sik | 1.<br>61<br>7 | 3.<br>6       | 8                 | 8                 | Alzheimer's & dementia :<br>the journal of the<br>Alzheimer's Association | 202<br>4          | Dec<br>1,<br>2025 |
| <a href="#">36993280</a> <br><a href="#">DOI</a>  | Accurate estimation of rare cell type fractions from tissue omics data via hierarchical deconvolu... | Huang,<br>Pengh<br>ui<br>...3<br>more...<br>Wang,<br>Jiebiao     | 0.<br>30<br>9 | 0             | 3                 | 1.5               | bioRxiv : the preprint<br>server for biology                              | 202<br>3          | Dec<br>1,<br>2025 |

|  |   |  |               |               |   |   |   |      |             |
|--|---|--|---------------|---------------|---|---|---|------|-------------|
| <a href="#">37577715</a> <br><a href="#">DOI</a>     | scMD: cell type deconvolution using single-cell DNA methylation references.     | Cai,<br>Manqi<br>...2<br>more...<br>Wang,<br>Jiebiao     | 0.<br>20<br>8 | 0             | 2 | 1 | bioRxiv : the preprint server for biology | 2023 | Dec 1, 2025 |
| <a href="#">39149243</a> <br><a href="#">DOI</a>     | BLEND: Probabilistic Cellular Deconvolution with Automated Reference Selection. | Huang,<br>Penghui<br>...2<br>more...<br>Wang,<br>Jiebiao | 0             | 0             | 0 | 0 | bioRxiv : the preprint server for biology | 2024 | Dec 1, 2025 |
| <a href="#">41112940</a> <br><a href="#">DOI</a>  | EMixed: Probabilistic Multi-Omics Cellular Deconvolution of Bulk Omics Data.    | Cai,<br>Manqi<br>...5<br>more...<br>Wang,<br>Jiebiao     | 0             | 0.<br>38<br>7 | 0 | 0 | Journal of data science : JDS             | 2025 | Dec 1, 2025 |





Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

# </> Repositories

Software repositories associated with this project.

C

Built on Dec 11, 2025

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

| Name    | Description | Tags | Last Commit | Statistics |         | Issues | PRs | Average   |        | Dependencies | Languages |
|---------|-------------|------|-------------|------------|---------|--------|-----|-----------|--------|--------------|-----------|
|         |             |      |             | Stars      | Commits |        |     | Issue Avg | PR Avg |              |           |
| 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.