



## (Core Project R03OD032630)

### ◎ Details

| Projects        | Name   | Award        | Publications   | Repositories   | Analytics    |
|-----------------|--|--------------|----------------|----------------|--------------|
| 1R03OD032630-01 | Methods to maximize the utility of common fund functional genomic data in multi-ethnic genetic studies | \$335,407.00 | 9 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>yea<br>r | Journal   | Publ<br>ishe<br>d | Upda<br>ted |
|--------------------------|--|---------------|---------|-----|-------------------|-------------------|-----------|-------------------|-------------|
| <a href="#">30643251</a> | Association studies of up to 1.2 million individuals yield new insights into the | Liu, Mengzhen | 65.06   | 0   | 1,390             | 231.667           | Nat Genet | 2019              | Nov 28,     |

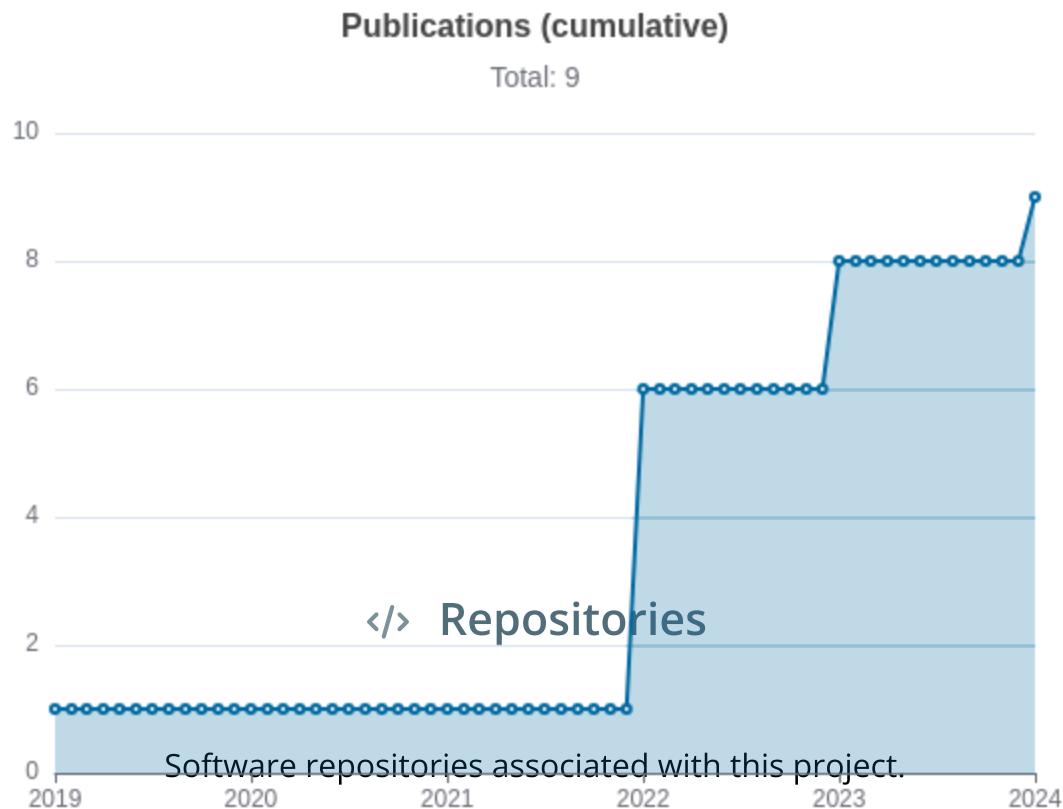
|                          |  |   |                |                |     |             |                   |   |
|--------------------------|--|---|----------------|----------------|-----|-------------|-------------------|---|
|                          | genetic etiology...  | ...123<br>more...<br>Vrieze,<br>Scott                                     | 5              |                |     |             |                   | 2025<br>(just<br>now)                       |
| <a href="#">36477530</a> | Genetic diversity fuels gene discovery for tobacco and alcohol use.                                  | Saunders,<br>Gretchen R<br>B<br>...217<br>more...<br>Vrieze,<br>Scott     | 30.<br>24<br>1 | 18.<br>28<br>8 | 323 | 107.<br>667 | Nature            | 2022<br>Nov<br>28,<br>2025<br>(just<br>now) |
| <a href="#">36750564</a> | Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict... | Khunsrirak<br>sakul,<br>Chachrit<br>...16<br>more...<br>Liu, Dajiang<br>J | 3.8<br>97      | 0              | 35  | 17.5        | Nat<br>Comm<br>un | 2023<br>Nov<br>28,<br>2025<br>(just<br>now) |
| <a href="#">36702996</a> | Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an... | Chen, Fang<br>...88<br>more...<br>Liu, Dajiang<br>J                       | 3.7<br>19      | 0              | 35  | 17.5        | Nat<br>Genet      | 2023<br>Nov<br>28,<br>2025<br>(just<br>now) |
| <a href="#">35672318</a> | Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing ... | Khunsrirak<br>sakul,<br>Chachrit  | 2.1<br>14      | 0              | 30  | 10          | Nat<br>Comm<br>un | 2022<br>Nov<br>28,<br>2025                  |



## Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 



| Name    | Tags | Last Commit | Avg Issue | Avg PR | Languages | License | Readme | Contributing | Dependencies |
|---------|------|-------------|-----------|--------|-----------|---------|--------|--------------|--------------|
| 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.

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 [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.

