





## 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 | 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>yea<br>r | Journal            | Publ<br>ishe<br>d | Upda<br>ted |
|---|--|-----------------|----------|----------|-------------------|-------------------|--------------------|-------------------|-------------|
| <a href="#">30643251</a> <br><a href="#">DOI</a>  | Association studies of up to 1.2 million individuals yield new insights into the | Mengzhen<br>Liu | 63<br>.3 | 17.<br>3 | 1,10<br>0         | 220               | Nature<br>Genetics | 2,01<br>9         | Oct<br>26,  |

|   |  |   |          |                |     |      |                          |           |   |                             |
|---|--|---|----------|----------------|-----|------|--------------------------|-----------|---|-----------------------------|
|   | genetic etiology...  | ...121<br>more...<br>Scott<br>Vrieze                                    | 4        |                |     |      |                          |           |   | 2024<br>(3<br>weeks<br>ago) |
| <a href="#">36477530</a> <br><a href="#">DOI</a>      | Genetic diversity fuels gene discovery for tobacco and alcohol use.                                  | Gretchen<br>R B<br>Saunders<br>...215<br>more...<br>Scott<br>Vrieze     | 23       | 18.<br>50<br>9 | 145 | 72.5 | Nature                   | 2,02<br>2 | Oct<br>26,<br>2024<br>(3<br>weeks<br>ago) |                             |
| <a href="#">36702996</a> <br><a href="#">DOI</a>      | Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an... | Fang Chen<br>...88<br>more...<br>Dajiang J<br>Liu                       | 4.<br>87 | 17.<br>3       | 19  | 19   | Nature<br>Genetics       | 2,02<br>3 | Oct<br>26,<br>2024<br>(3<br>weeks<br>ago) |                             |
| <a href="#">36750564</a> <br><a href="#">DOI</a>  | Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict... | Chachrit<br>Khunsrirak<br>sakul<br>...16<br>more...<br>Dajiang J<br>Liu | 3.<br>55 | 4.8<br>87      | 14  | 14   | Nature<br>Communications | 2,02<br>3 | Oct<br>26,<br>2024<br>(3<br>weeks<br>ago) |                             |
| <a href="#">35672318</a> <br><a href="#">DOI</a>  | Integrating 3D genomic and epigenomic data to enhance target gene discovery                          | Chachrit<br>Khunsrirak  | 1.<br>68 | 4.8<br>87      | 15  | 7.5  | Nature<br>Communic       | 2,02<br>2 | Oct<br>26,                                |                             |

|   |   |   |          |           |    |     |  |                                   |           |   |
|---|---|---|----------|-----------|----|-----|--|-----------------------------------|-----------|---|
|   | and drug repurposing ...  | sakul<br>...12<br>more...<br>Dajiang J<br>Liu                       |          |           |    |     |  | ations                            |           | 2024<br>(3<br>weeks<br>ago)               |
| <a href="#">35927319</a> <br><a href="#">DOI</a>      | Rare genetic variants explain missing heritability in smoking.                        | Seon-<br>Kyeong<br>Jang<br>...88<br>more...<br>Scott<br>Vrieze      | 1.<br>52 | 6.0<br>97 | 13 | 6.5 |  | Nature<br>Human<br>Behaviour      | 2,02<br>2 | Oct<br>26,<br>2024<br>(3<br>weeks<br>ago) |
| <a href="#">35833142</a> <br><a href="#">DOI</a>     | Construction and Application of Polygenic Risk Scores in Autoimmune Diseases.         | Chachrit<br>Khunsrirak<br>sakul<br>...4 more...<br>Dajiang J<br>Liu | 0.<br>91 | 1.8<br>68 | 7  | 3.5 |  | Frontiers<br>in<br>Immunolo<br>gy | 2,02<br>2 | Oct<br>26,<br>2024<br>(3<br>weeks<br>ago) |
| <a href="#">35178743</a> <br><a href="#">DOI</a>  | Assessing reproducibility of high-throughput experiments in the case of missing data. | Roopali<br>Singh<br>...1 more...<br>Qunhua Li                       | 0.<br>48 | 1.3<br>48 | 3  | 1.5 |  | Statistics<br>in<br>Medicine      | 2,02<br>2 | Oct<br>26,<br>2024<br>(3<br>weeks<br>ago) |

Notes

RCR [Relative Citation Ratio](#)

SJR [Scimago Journal Rank](#)



| 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

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.