

# **L** Core Project R03OD032630

### O 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 |
|------------|--|------------------|-----------|-----|-------------------|-------------------|--------------|-------------------|-------------|
| 30643251 🗹 | Association studies of up to 1.2 million individuals yield new insights into the | Liu,<br>Mengzhen | 64.<br>91 | 0   | 1,34<br>7         | 224.<br>5         | Nat<br>Genet | 2019              | Sep<br>3,   |

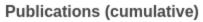
|                     | genetic etiology  | 123<br>more<br>Vrieze,<br>Scott                                  | 7              |                |     |            |                   |      | 2025<br>(just<br>now)              |
|---------------------|---|--|----------------|----------------|-----|------------|-------------------|------|------------------------------------|
| 36477530 🗗          | Genetic diversity fuels gene discovery for tobacco and alcohol use.                               | Saunders,<br>Gretchen R<br>B<br>217<br>more<br>Vrieze,<br>Scott  | 28.<br>38<br>7 | 18.<br>28<br>8 | 286 | 95.3<br>33 | Nature            | 2022 | Sep<br>3,<br>2025<br>(just<br>now) |
| 36702996 ☑<br>DOI ☑ | Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an | Chen, Fang<br>88 more<br>Liu, Dajiang<br>J                       | 4.0<br>51      | 0              | 34  | 17         | Nat<br>Genet      | 2023 | Sep<br>3,<br>2025<br>(just<br>now) |
| 36750564 🗳          | Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict | Khunsrirak<br>sakul,<br>Chachrit<br>16 more<br>Liu, Dajiang<br>J | 3.6<br>92      | 0              | 30  | 15         | Nat<br>Comm<br>un | 2023 | Sep<br>3,<br>2025<br>(just<br>now) |
| 35672318 🗹          | Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing  | Khunsrirak<br>sakul,<br>Chachrit<br>12 more                      | 2.0<br>95      | 0              | 28  | 9.33<br>3  | Nat<br>Comm<br>un | 2022 | Sep<br>3,<br>2025                  |

|                                   |  | Liu, Dajiang<br>J   |           |   |    |           |                      |      | (just<br>now)                      |
|-----------------------------------|--|---|-----------|---|----|-----------|----------------------|------|------------------------------------|
| 35927319 ♂<br>DOI ♂               | Rare genetic variants explain missing heritability in smoking.                                 | Jang, Seon-<br>Kyeong<br>88 more<br>Vrieze,<br>Scott            | 1.4<br>53 | 0 | 17 | 5.66<br>7 | Nat<br>Hum<br>Behav  | 2022 | Sep<br>3,<br>2025<br>(just<br>now) |
| 35833142 <b>亿</b><br>DOI <b>亿</b> | Construction and Application of Polygenic<br>Risk Scores in Autoimmune Diseases.               | Khunsrirak<br>sakul,<br>Chachrit<br>4 more<br>Liu, Dajiang<br>J | 1.0<br>37 | 0 | 12 | 4         | Front<br>Immun<br>ol | 2022 | Sep<br>3,<br>2025<br>(just<br>now) |
| 35178743 🗗                        | Assessing reproducibility of high-<br>throughput experiments in the case of<br>missing data.   | Singh,<br>Roopali<br>1 more<br>Li, Qunhua                       | 0.4<br>91 | 0 | 5  | 1.66<br>7 | Stat<br>Med          | 2022 | Sep<br>3,<br>2025<br>(just<br>now) |
| 38918381 ☑<br>DOI ☑               | Dissecting heritability, environmental risk, and air pollution causal effects using > 50 milli | McGuire,<br>Daniel<br>8 more<br>Jiang, Bibo                     | 0         | 0 | 1  | 1         | Nat<br>Comm<br>un    | 2024 | Sep<br>3,<br>2025<br>(just<br>now) |

### Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank





| Name | Description | Stars | Watchers | Forks  | Issues | PRs | Commits | Contrib. |
|------|-------------|-------|----------|--------|--------|-----|---------|----------|
|      |             |       | N        | o 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 2.

New Users <u>Users who visited the website for the first time</u> **.** 

Engaged Sessions <u>Visits that had significant interaction</u> **?**.

"Top" metrics are measured by number of engaged sessions.