

# **▶** 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 | 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 | Updat<br>ed     |
|------------|--|-----------------|----------|----------|-------------------|-------------------|--------------------|-------------------|-----------------|
| 30643251 🗹 | Association studies of up to 1.2 million individuals yield new insights into the | Mengzhen<br>Liu | 62<br>.9 | 17.<br>3 | 1,08<br>5         | 217               | Nature<br>Genetics | 2,01<br>9         | Sep 28,<br>2024 |

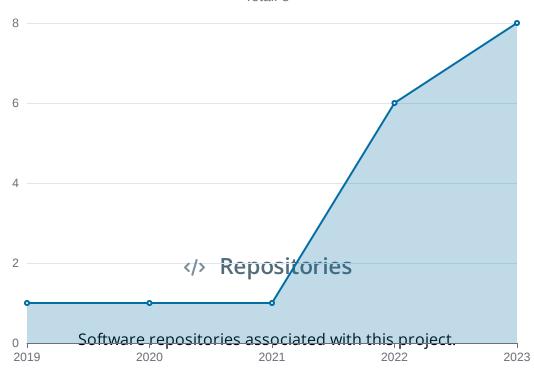
|                                   | genetic etiology   | 121<br>more<br>Scott<br>Vrieze                                    | 4             |                |     |          |                              |           | (3<br>weeks<br>ago)                       |
|-----------------------------------|--|---|---------------|----------------|-----|----------|------------------------------|-----------|---|
| 36477530 <b>乙</b><br>DOI <b>乙</b> | Genetic diversity fuels gene discovery for tobacco and alcohol use.                                      | Gretchen<br>R B<br>Saunders<br>215<br>more<br>Scott<br>Vrieze     | 21<br>.1<br>9 | 18.<br>50<br>9 | 129 | 64.<br>5 | Nature                       | 2,02<br>2 | Sep 28,<br>2024<br>(3<br>weeks<br>ago)    |
| 36702996 🗗<br>DOI 🗗               | Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an        | Fang Chen<br>88<br>more<br>Dajiang J<br>Liu                       | 5.<br>2       | 17.<br>3       | 18  | 18       | Nature<br>Genetics           | 2,02<br>3 | Oct 18,<br>2024<br>(9<br>minute<br>s ago) |
| 36750564 ♂<br>DOI ♂               | Multi-ancestry and multi-trait genome-<br>wide association meta-analyses inform<br>clinical risk predict | Chachrit<br>Khunsrira<br>ksakul<br>16<br>more<br>Dajiang J<br>Liu | 3.<br>97      | 4.8<br>87      | 14  | 14       | Nature<br>Communic<br>ations | 2,02<br>3 | Sep 28,<br>2024<br>(3<br>weeks<br>ago)    |
| 35672318 🗗                        | Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing         | Chachrit<br>Khunsrira<br>ksakul                                   | 1.<br>71      | 4.8<br>87      | 15  | 7.5      | Nature<br>Communic<br>ations | 2,02<br>2 | Sep 28,<br>2024<br>(3                     |

|                                   |   | 12<br>more<br>Dajiang J<br>Liu                                |          |           |    |     |                                   |           | weeks<br>ago)                          |
|-----------------------------------|---|---|----------|-----------|----|-----|-----------------------------------|-----------|--|
| 35927319 <b>♂</b><br>DOI <b>♂</b> | Rare genetic variants explain missing heritability in smoking.                | Seon-<br>Kyeong<br>Jang<br>88<br>more<br>Scott<br>Vrieze      | 1.<br>46 | 6.0<br>97 | 12 | 6   | Nature<br>Human<br>Behaviour      | 2,02<br>2 | Sep 28,<br>2024<br>(3<br>weeks<br>ago) |
| 35833142 ☑<br>DOI ☑               | Construction and Application of Polygenic Risk Scores in Autoimmune Diseases. | Chachrit<br>Khunsrira<br>ksakul<br>4 more<br>Dajiang J<br>Liu | 0.<br>93 | 1.8<br>68 | 7  | 3.5 | Frontiers<br>in<br>Immunolo<br>gy | 2,02<br>2 | Sep 28,<br>2024<br>(3<br>weeks<br>ago) |
| <u>35178743</u> <b>∠</b>          | Assessing reproducibility of high-<br>throughput experiments in the case of   | Roopali<br>Singh  | 0.       | 1.3       | 2  | 1   | Statistics<br>in                  | 2,02      | Sep 28,<br>2024<br>(3                  |

# Notes

#### **Cumulative Publications**





| 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   |         |        |              |              |

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 <u>Distinct users who visited the website</u> **?**.

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

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