

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 R | SJR | Cita tion s | Cit./ yea r | Journal | Publ ishe d | Upda ted |
|------------|--|------------------|-----------|-----|-------------------|-------------------|--------------|-------------------|-------------|
| 30643251 🗹 | Association studies of up to 1.2 million individuals yield new insights into the | Liu, Mengzhen | 64. 91 | 0 | 1,34 7 | 224. 5 | Nat Genet | 2019 | Sep 21, |

| | genetic etiology | 123 more Vrieze, Scott | 7 | | | | | | 2025 (just now) |
|-----------------------------------|---|---|----------------|----------------|-----|------------|-------------------|------|-------------------------------------|
| 36477530 乙 DOI 乙 | Genetic diversity fuels gene discovery for tobacco and alcohol use. | Saunders, Gretchen R B217 more Vrieze, Scott | 28. 38 7 | 18. 28 8 | 286 | 95.3 33 | Nature | 2022 | Sep 21, 2025 (just now) |
| 36702996 🗗 DOI 🗗 | Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an | Chen, Fang 88 more Liu, Dajiang J | 4.0 51 | 0 | 34 | 17 | Nat Genet | 2023 | Sep 21, 2025 (just now) |
| 36750564 乙 | Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict | Khunsrirak sakul, Chachrit 16 more Liu, Dajiang J | 3.6 92 | 0 | 30 | 15 | Nat Comm un | 2023 | Sep 21, 2025 (just now) |
| 35672318 🗗 | Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing | Khunsrirak sakul, Chachrit | 2.0 95 | 0 | 28 | 9.33 3 | Nat Comm un | 2022 | Sep 21, 2025 |

| | | 12 more Liu, Dajiang J | | | | | | | (just now) |
|-----------------------------------|--|---|-----------|---|----|-----------|----------------------|------|-------------------------------------|
| 35927319 ♂ DOI ♂ | Rare genetic variants explain missing heritability in smoking. | Jang, Seon- Kyeong 88 more Vrieze, Scott | 1.4 53 | 0 | 17 | 5.66 7 | Nat Hum Behav | 2022 | Sep 21, 2025 (just now) |
| 35833142 ☑ DOI ☑ | Construction and Application of Polygenic Risk Scores in Autoimmune Diseases. | Khunsrirak sakul, Chachrit 4 more Liu, Dajiang J | 1.0 37 | 0 | 12 | 4 | Front Immun ol | 2022 | Sep 21, 2025 (just now) |
| 35178743 🗗 | Assessing reproducibility of high- throughput experiments in the case of missing data. | Singh, Roopali 1 more Li, Qunhua | 0.4 91 | 0 | 5 | 1.66 7 | Stat Med | 2022 | Sep 21, 2025 (just now) |
| 38918381 ♂ DOI ♂ | Dissecting heritability, environmental risk, and air pollution causal effects using > 50 milli | McGuire, Daniel 8 more Jiang, Bibo | 0 | 0 | 1 | 1 | Nat Comm un | 2024 | Sep 21, 2025 (just now) |

Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank

Publications (cumulative)



| Name | Description | Stars | Watchers | Forks | Issues | PRs | Commits | Contrib. |
|------|-------------|-------|----------|--------|--------|-----|---------|----------|
| | | | N | o 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 <u>Distinct users who visited the website</u> 2.

New Users Users who visited the website for the first time **2**.

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

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