



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 R | SJR | Cita tion s | Cit./ yea r | Journal | Publ ishe d | Upda ted |
|---|--|------------------|-----------|-----|-------------------|-------------------|--------------|-------------------|-------------|
| 30643251  DOI  | Association studies of up to 1.2 million individuals yield new insights into the | Liu, Mengzhen | 65. 06 | 0 | 1,36 8 | 228 | Nat Genet | 2019 | Oct 2, |

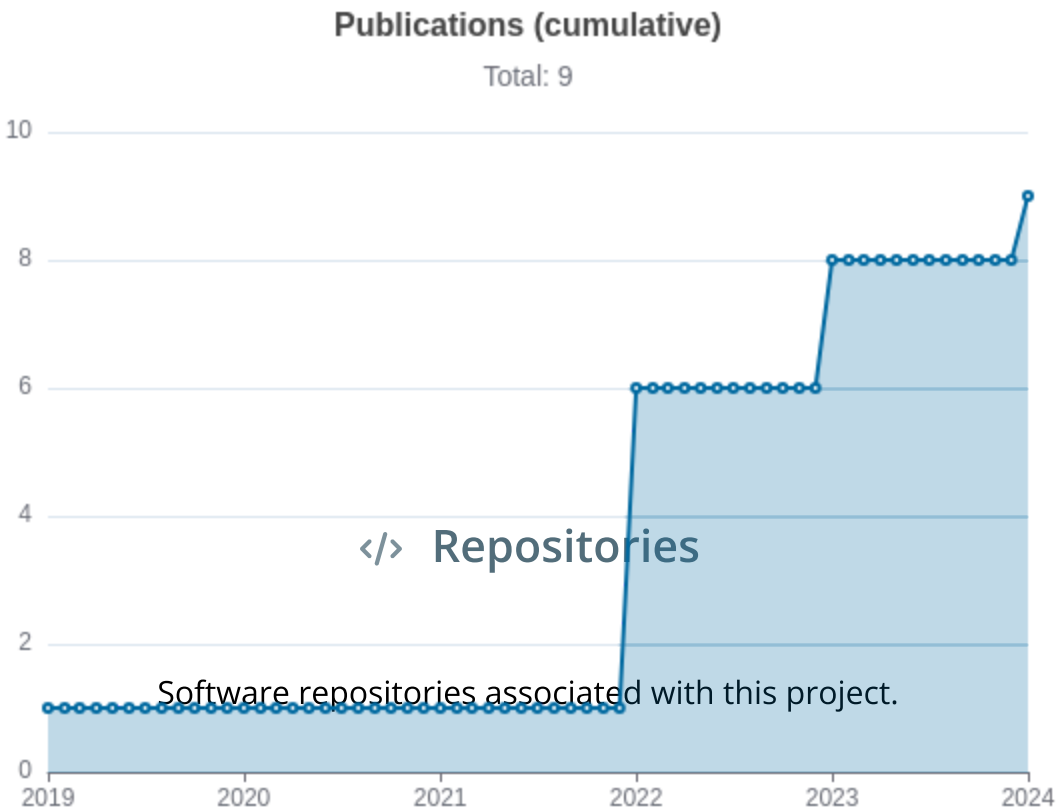
| | | | | | | | | | | |
|--|--|--|----------------|----------------|-----|-----------|-------------------|------|------------------------------------|-----------------------|
| | genetic etiology... | ...123 more... Vrieze, Scott | 1 | | | | | | | 2025 (just now) |
| 36477530 DOI ↗ | Genetic diversity fuels gene discovery for tobacco and alcohol use. | Saunders, Gretchen R B ...217 more... Vrieze, Scott | 29. 55 5 | 18. 28 8 | 306 | 102 | Nature | 2022 | Oct 2, 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 | 3.9 55 | 0 | 35 | 17.5 | Nat Genet | 2023 | Oct 2, 2025 (just now) | |
| 36750564 DOI ↗ | Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict... | Khunsrirak sakul, Chachrit ...16 more... Liu, Dajiang J | 3.7 01 | 0 | 31 | 15.5 | Nat Comm un | 2023 | Oct 2, 2025 (just now) | |
| 35672318 DOI ↗ | Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing ... | Khunsrirak sakul, Chachrit ...12 more... | 2.0 36 | 0 | 28 | 9.33 3 | Nat Comm un | 2022 | Oct 2, 2025 | |

| | | | | | | | | | | |
|--|---|--|-------|---|----|-------|---------------|------|------------------------|------------|
| | | Liu, Dajiang J | | | | | | | | (just now) |
| 35927319 DOI ↗ | Rare genetic variants explain missing heritability in smoking. | Jang, Seon-Kyeong ...88 more... Vrieze, Scott | 1.411 | 0 | 17 | 5.667 | Nat Hum Behav | 2022 | Oct 2, 2025 (just now) | |
| 35833142 DOI ↗ | Construction and Application of Polygenic Risk Scores in Autoimmune Diseases. | Khunsrirak sakul, Chachrit ...4 more... Liu, Dajiang J | 1.095 | 0 | 13 | 4.333 | Front Immunol | 2022 | Oct 2, 2025 (just now) | |
| 35178743 DOI ↗ | Assessing reproducibility of high-throughput experiments in the case of missing data. | Singh, Roopali ...1 more... Li, Qunhua | 0.477 | 0 | 5 | 1.667 | Stat Med | 2022 | Oct 2, 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 Commun | 2024 | Oct 2, 2025 (just now) | |

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.