



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

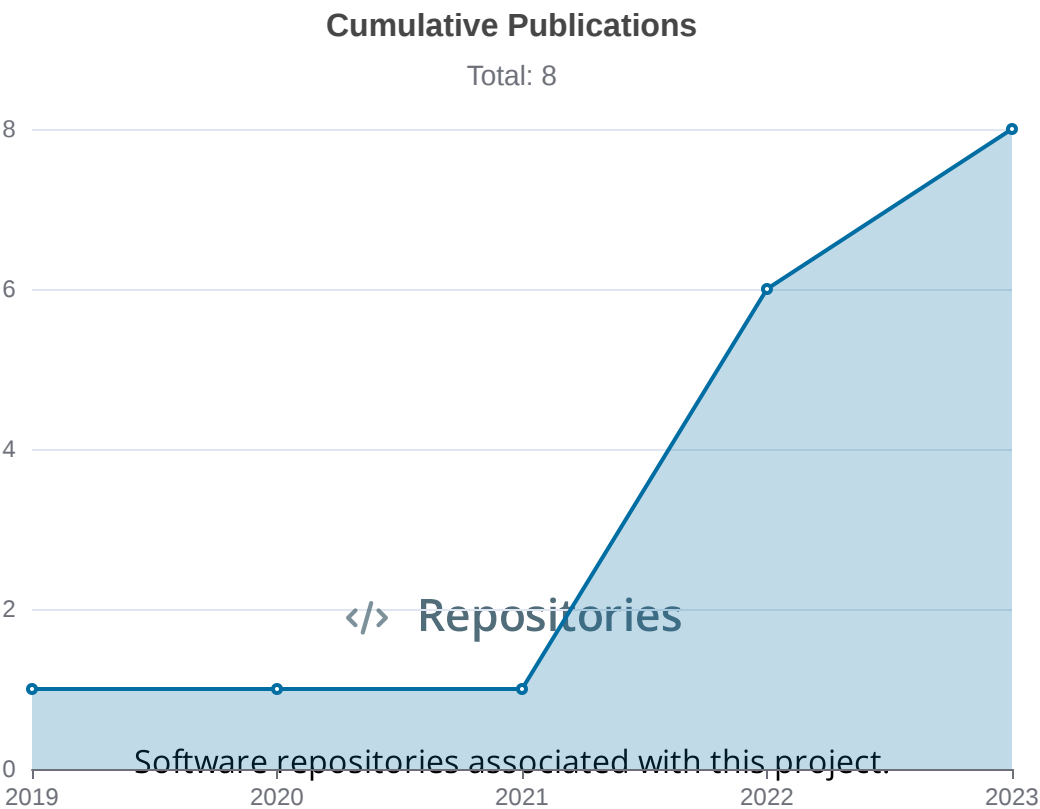
Published works associated with this project.

| ID | Title | Authors | RC 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 | Mengzhen Liu | 62 .2 | 17. 3 | 1,06 8 | 213. 6 | Nature Genetics | 2,01 9 | Sep 1, 2024 |

| | | | | | | | | | | |
|---|--|---|---------------|----------------|-----|------|------------------------------|-----------|--|---------------------|
| | genetic etiology... | ...121 more... Scott Vrieze | 5 | | | | | | | (3 weeks ago) |
| 36477530  DOI  | Genetic diversity fuels gene discovery for tobacco and alcohol use. | Gretchen R B Saunders ...215 more... Scott Vrieze | 21 .3 7 | 18. 50 9 | 125 | 62.5 | Nature | 2,02 2 | Sep 1, 2024 (3 weeks ago) | |
| 36702996  DOI  | Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an... | Fang Chen ...88 more... Dajiang J Liu | 5. 62 | 17. 3 | 18 | 18 | Nature Genetics | 2,02 3 | Sep 1, 2024 (3 weeks ago) | |
| 36750564  DOI  | Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict... | Chachrit Khunsrirak sakul ...16 more... Dajiang J Liu | 4. 29 | 4.8 87 | 14 | 14 | Nature Communic ations | 2,02 3 | Sep 13, 2024 (1 week ago) | |
| 35672318  DOI  | Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing ... | Chachrit Khunsrirak sakul | 1. 66 | 4.8 87 | 14 | 7 | Nature Communic ations | 2,02 2 | Sep 1, 2024 (3 | |

| | | | | | | | | | | |
|---|---|---|----------|-----------|----|-----|-----------------------------------|-----------|---------------------------------------|---------------|
| | | ...12 more... Dajiang J Liu | | | | | | | | weeks ago) |
| 35927319  DOI  | Rare genetic variants explain missing heritability in smoking. | Seon- Kyeong Jang ...88 more... Scott Vrieze | 1. 51 | 6.0 97 | 12 | 6 | Nature Human Behaviour | 2,02 2 | Sep 7, 2024 (2 weeks ago) | |
| 35833142  DOI  | Construction and Application of Polygenic Risk Scores in Autoimmune Diseases. | Chachrit Khunsrirak sakul ...4 more... Dajiang J Liu | 0. 97 | 1.8 68 | 7 | 3.5 | Frontiers in Immunolo gy | 2,02 2 | Sep 1, 2024 (3 weeks ago) | |
| 35178743  DOI  | Assessing reproducibility of high-throughput experiments in the case of missing data. | Roopali Singh ...1 more... Qunhua Li | 0. 32 | 1.3 48 | 2 | 1 | Statistics in Medicine | 2,02 2 | Sep 1, 2024 (3 weeks ago) | |

Notes



| 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.