

▶ 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 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 | Mengzhen Liu | 62 .9 | 17. 3 | 1,08 5 | 217 | Nature Genetics | 2,01 9 | Sep 28, |

| | genetic etiology | 121 more Scott Vrieze | 4 | | | | | | 2024 (3 weeks ago) |
|-----------------------------------|--|---|---------------|----------------|-----|------|------------------------------|-----------|---|
| 36477530 ♂ DOI ♂ | Genetic diversity fuels gene discovery for tobacco and alcohol use. | Gretchen R B Saunders 215 more Scott Vrieze | 21 .1 9 | 18. 50 9 | 129 | 64.5 | Nature | 2,02 2 | Sep 28, 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. 2 | 17. 3 | 18 | 18 | Nature Genetics | 2,02 3 | Sep 28, 2024 (3 weeks ago) |
| <u>36750564</u> ♂ <u>DOI</u> ♂ | Multi-ancestry and multi-trait genome- wide association meta-analyses inform clinical risk predict | Chachrit Khunsrirak sakul 16 more Dajiang J Liu | 3. 97 | 4.8 87 | 14 | 14 | Nature Communic ations | 2,02 3 | Sep 28, 2024 (3 weeks ago) |
| 35672318 🗹 | Integrating 3D genomic and epigenomic data to enhance target gene discovery | Chachrit Khunsrirak | 1. 71 | 4.8 87 | 15 | 7.5 | Nature Communic | 2,02 2 | Sep 28, |

| | and drug repurposing | sakul 12 more Dajiang J Liu | | | | | ations | | 2024 (3 weeks ago) |
|---------------------|--|---|----------|-----------|----|-----|-----------------------------------|-----------|---|
| 35927319 🗗 | Rare genetic variants explain missing heritability in smoking. | Seon- Kyeong Jang 88 more Scott Vrieze | 1. 46 | 6.0 97 | 12 | 6 | Nature Human Behaviour | 2,02 2 | Sep 28, 2024 (3 weeks ago) |
| 35833142 乙 | Construction and Application of Polygenic Risk Scores in Autoimmune Diseases. | Chachrit Khunsrirak sakul 4 more Dajiang J Liu | 0. 93 | 1.8 68 | 7 | 3.5 | Frontiers in Immunolo gy | 2,02 2 | Sep 28, 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. 3 | 1.3 48 | 2 | 1 | Statistics in Medicine | 2,02 2 | Sep 28, 2024 (3 weeks ago) |

Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank





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

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

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