



Core Project R03OD032630









Overview











High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD032630-01	Methods to maximize the utility of common fund functional genomic data in multi-ethnic genetic studies	\$335K	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 genetic etiology...	Liu, Mengzhen ...123 more... Vrieze, Scott	65. 25 7	16. 58 6	1,43 1	238. 5	Nature genetics	2019	Dec 28, 2025
36477530  DOI 	Genetic diversity fuels gene discovery for tobacco and alcohol use.	Saunders, Gretchen R B ...217 more... Vrieze, Scott	30. 79 7	18. 28 8	346	115. 333	Nature	2022	Dec 28, 2025
36750564  DOI 	Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict...	Khunsriraks akul, Chachrit ...16 more... Liu, Dajiang J	4.2 71	4.7 61	41	20.5	Nature communica tions	2023	Dec 28, 2025

36702996  DOI 	Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an...	Chen, Fang ...88 more... Liu, Dajiang J	3.8 35	16. 58 6	39	19.5	Nature genetics	2023	Dec 28, 2025
35672318  DOI 	Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing ...	Khunsriraks akul, Chachrit ...12 more... Liu, Dajiang J	2.2 23	4.7 61	33	11	Nature communica tions	2022	Dec 28, 2025
35927319  DOI 	Rare genetic variants explain missing heritability in smoking.	Jang, Seon- Kyeong ...88 more... Vrieze, Scott	1.4 93	5.5 37	20	6.66 7	Nature human behaviour	2022	Dec 28, 2025
35833142  DOI 	Construction and Application of Polygenic Risk Scores in Autoimmune Diseases.	Khunsriraks akul, Chachrit ...4 more... Liu, Dajiang J	1.0 08	1.9 41	13	4.33 3	Frontiers in immunolog y	2022	Dec 28, 2025
38918381  DOI 	Dissecting heritability, environmental risk, and air pollution causal effects using > 50 milli...	McGuire, Daniel ...8 more... Jiang, Bibo	0.9 02	4.7 61	2	2	Nature communica tions	2024	Dec 28, 2025

[35178743](#) [DOI](#)

Assessing reproducibility of high-throughput experiments in the case of missing data.

Singh,
Roopali
...1 more...
Li, Qunhua

0.5
1

1.2
68

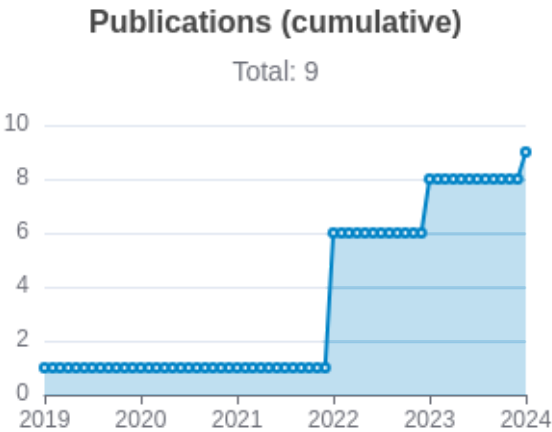
6

2

Statistics in
medicine

2022

Dec
28,
2025



Notes

- RCR [Relative Citation Ratio](#)
- SJR [Scimago Journal Rank](#)

</> Repositories

Software repositories associated with this project.

Name	Description	Tags	Last Commit	Stars	Forks	Watchers	Commits	Issues	PRs	Issue Avg	PR Avg	Readme	Contributing	Code of Con.	License	Contribs	Languages

Built on Jan 30, 2026

Developed with support from NIH Award [U54 OD036472](#)

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.
- Issue/PR Avg 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

Website metrics 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.