



## (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
<a href="#">30643251</a>	Association studies of up to 1.2 million individuals yield new insights into the	Liu, Mengzhen	65. 06	0	1,39 0	231. 667	Nat Genet	2019	Nov 20,

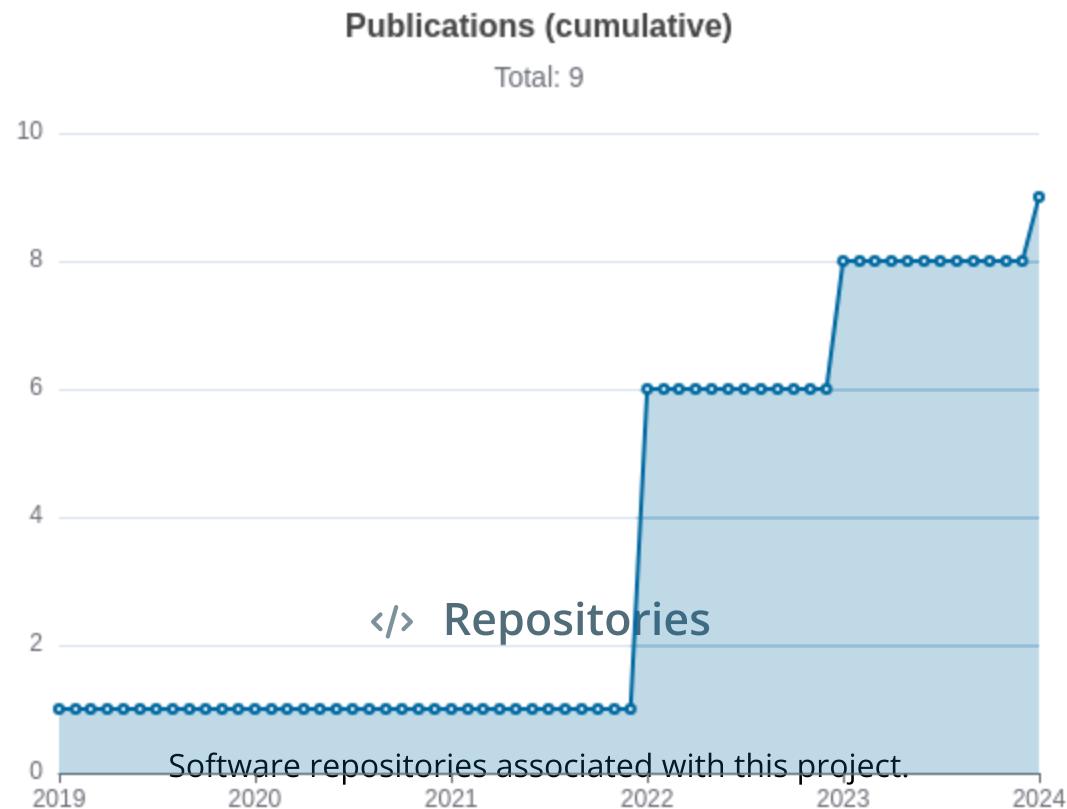
	genetic etiology...	...123 more... Vrieze, Scott	5					2025 (just now)
<a href="#">36477530</a>	Genetic diversity fuels gene discovery for tobacco and alcohol use.	Saunders, Gretchen R B ...217 more... Vrieze, Scott	30. 24 1	18. 28 8	323	107. 667	Nature	2022 Nov 20, 2025 (just now)
<a href="#">36750564</a>	Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict...	Khunsrirak sakul, Chachrit ...16 more... Liu, Dajiang J	3.8 97	0	35	17.5	Nat Comm un	2023 Nov 20, 2025 (just now)
<a href="#">36702996</a>	Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an...	Chen, Fang ...88 more... Liu, Dajiang J	3.7 19	0	35	17.5	Nat Genet	2023 Nov 20, 2025 (just now)
<a href="#">35672318</a>	Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing ...	Khunsrirak sakul, Chachrit	2.1 14	0	30	10	Nat Comm un	2022 Nov 20, 2025



## Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 



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.

