





## 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	Updat ed
<a href="#">30643251</a>  <a href="#">DOI</a> 	Association studies of up to 1.2 million individuals yield new insights into the	Mengzhen Liu	63. 31	0	1,12 9	225. 8	Nat Genet	2019	Dec 28,

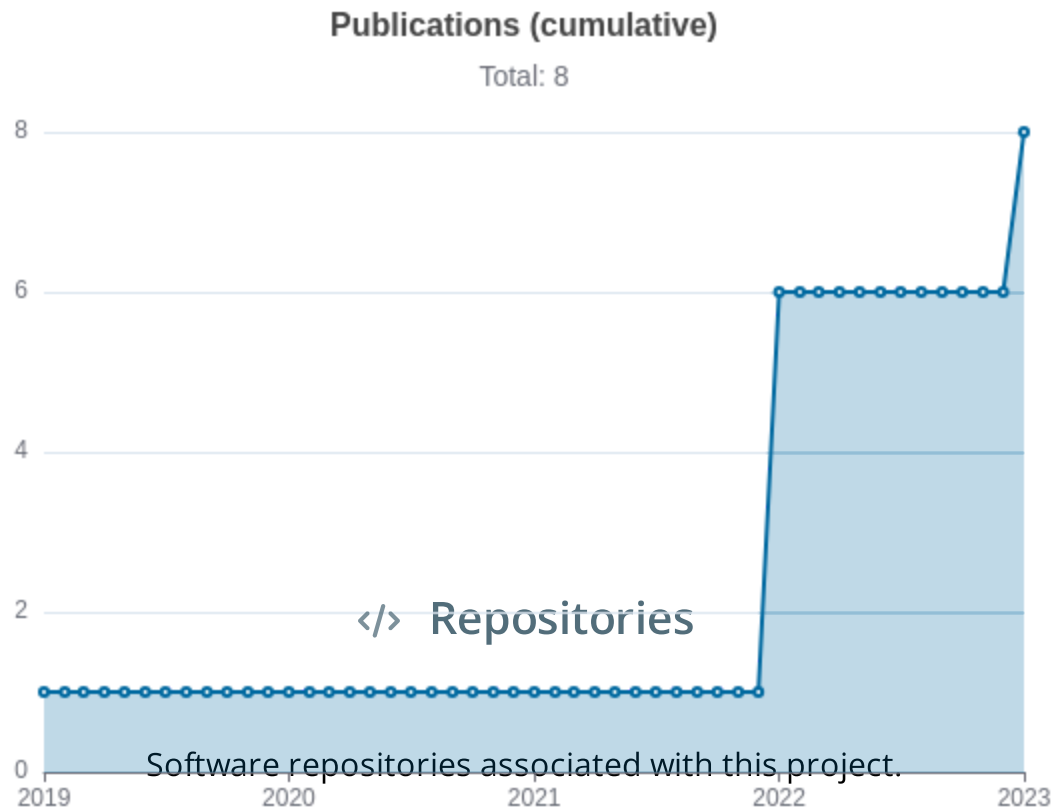
	genetic etiology...	...121 more... Scott Vrieze								2024 (2 weeks ago)
<a href="#">36477530</a>  <a href="#">DOI</a> 	Genetic diversity fuels gene discovery for tobacco and alcohol use.	Gretchen R B Saunders ...215 more... Scott Vrieze	23. 29	18. 50 9	159	79.5	Nature	2022	Dec 29, 2024 (2 weeks ago)	
<a href="#">36702996</a>  <a href="#">DOI</a> 	Multi-ancestry transcriptome-wide association analyses yield insights into tobacco use biology an...	Fang Chen ...88 more... Dajiang J Liu	5.0 4	0	22	22	Nat Genet	2023	Dec 28, 2024 (2 weeks ago)	
<a href="#">36750564</a>  <a href="#">DOI</a> 	Multi-ancestry and multi-trait genome-wide association meta-analyses inform clinical risk predict...	Chachrit Khunsrirak sakul ...16 more... Dajiang J Liu	3.4 8	0	15	15	Nat Comm un	2023	Dec 28, 2024 (2 weeks ago)	
<a href="#">35672318</a>  <a href="#">DOI</a> 	Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing ...	Chachrit Khunsrirak sakul ...12 more...	1.8 4	0	16	8	Nat Comm un	2022	Dec 29, 2024 (2	

		Dajiang J Liu								weeks ago)
<a href="#">35927319</a>  <a href="#">DOI</a> 	Rare genetic variants explain missing heritability in smoking.	Seon-Kyeong Jang ...88 more... Scott Vrieze	1.4 3	0	13	6.5	Nat Hum Behav	2022	Dec 29, 2024 (2 weeks ago)	
<a href="#">35833142</a>  <a href="#">DOI</a> 	Construction and Application of Polygenic Risk Scores in Autoimmune Diseases.	Chachrit Khunsrirak sakul ...4 more... Dajiang J Liu	0.8 6	0	7	3.5	Front Immunol	2022	Dec 29, 2024 (2 weeks ago)	
<a href="#">35178743</a>  <a href="#">DOI</a> 	Assessing reproducibility of high-throughput experiments in the case of missing data.	Roopali Singh ...1 more... Qunhua Li	0.4 5	0	3	1.5	Stat Med	2022	Dec 28, 2024 (2 weeks ago)	

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.