




## 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>  <a href="#">DOI</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 29,

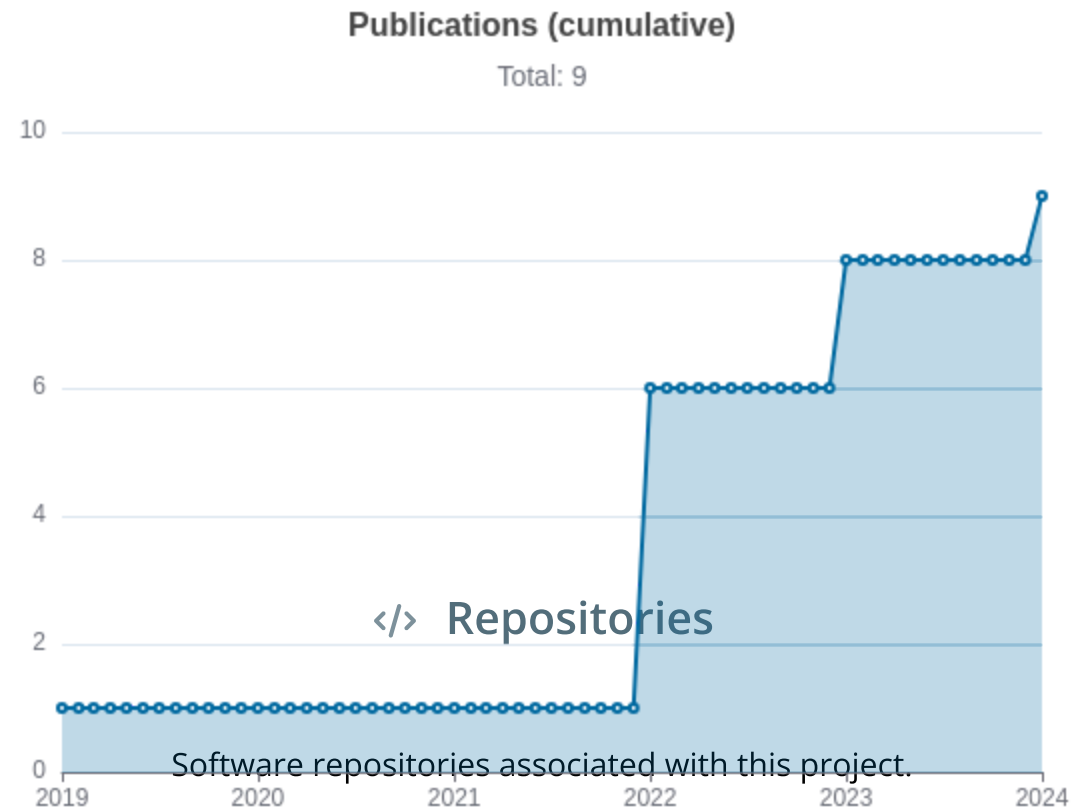
	genetic etiology...	...123 more... Vrieze, Scott	5							2025 (just now)
<a href="#">36477530</a>  <a href="#">DOI</a> 	Genetic diversity fuels gene discovery for tobacco and alcohol use.	Saunders, Gretchen R B ...217 more... Vrieze, Scott	30. 24 1	18. 28 8	323	107. 667	Nature	2022		Nov 29, 2025 (just now)
<a href="#">36750564</a>  <a href="#">DOI</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 29, 2025 (just now)
<a href="#">36702996</a>  <a href="#">DOI</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 29, 2025 (just now)
<a href="#">35672318</a>  <a href="#">DOI</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 29, 2025

		...12 more... Liu, Dajiang J								(just now)
<a href="#">35927319</a>  <a href="#">DOI</a> 	Rare genetic variants explain missing heritability in smoking.	Jang, Seon-Kyeong ...88 more... Vrieze, Scott	1.4 27	0	18	6	Nat Hum Behav	2022	Nov 29, 2025 (just now)	
<a href="#">35833142</a>  <a href="#">DOI</a> 	Construction and Application of Polygenic Risk Scores in Autoimmune Diseases.	Khunsrirak sakul, Chachrit ...4 more... Liu, Dajiang J	1.0 62	0	13	4.33 3	Front Immun ol	2022	Nov 29, 2025 (just now)	
<a href="#">38918381</a>  <a href="#">DOI</a> 	Dissecting heritability, environmental risk, and air pollution causal effects using &gt; 50 milli...	McGuire, Daniel ...8 more... Jiang, Bibo	1.0 43	0	2	2	Nat Comm un	2024	Nov 29, 2025 (just now)	
<a href="#">35178743</a>  <a href="#">DOI</a> 	Assessing reproducibility of high-throughput experiments in the case of missing data.	Singh, Roopali ...1 more... Li, Qunhua	0.4 59	0	5	1.66 7	Stat Med	2022	Nov 29, 2025 (just now)	

## 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 <code>main</code> /default branch is considered for metrics like # of commits.	
# of dependencies is totaled from all manifests in repo, direct and transitive, e.g. <code>package.json</code> + <code>package-lock.json</code> .	

 Analytics

Traffic metrics of websites associated with this project.

Notes

Active Users	<a href="#">Distinct users who visited the website</a>  .
New Users	<a href="#">Users who visited the website for the first time</a>  .
Engaged Sessions	<a href="#">Visits that had significant interaction</a>  .

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

