



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

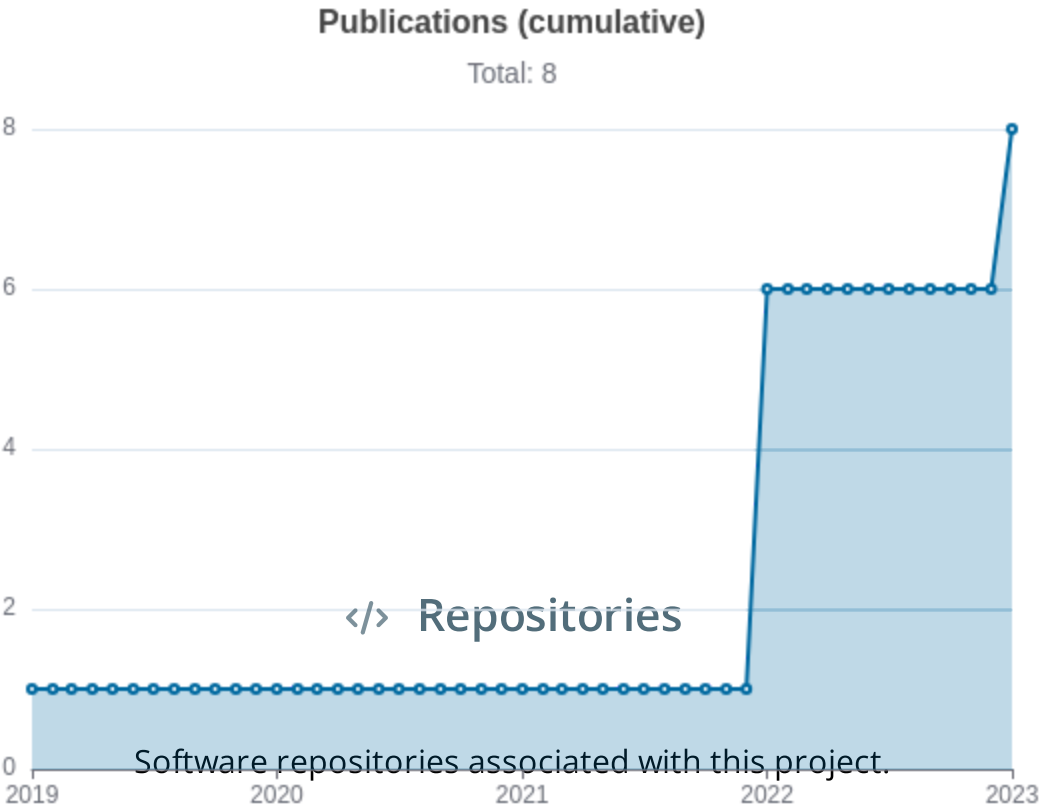
Published works associated with this project.

ID	Title	Authors	RC R	SJR	Cita tion s	Cit./ yea r	Journal	Publ ishe d	Update d
30643251  DOI 	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, 2024

	genetic etiology...	...121 more... Scott Vrieze								(52 minute s ago)
36477530  DOI 	Genetic diversity fuels gene discovery for tobacco and alcohol use.	Gretchen R B Saunders ...215 more... Scott Vrieze	23. 59	18. 50 9	155	77.5	Nature	2022	Dec 1, 2024 (1 month ago)	
36702996  DOI 	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 (49 minute s ago)	
36750564  DOI 	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 (46 minute s ago)	
35672318  DOI 	Integrating 3D genomic and epigenomic data to enhance target gene discovery and drug repurposing ...	Chachrit Khunsrirak sakul	1.8 4	0	16	8	Nat Comm un	2022	Dec 29, 2024 (14	

		...12 more... Dajiang J Liu								minute s ago)
35927319  DOI 	Rare genetic variants explain missing heritability in smoking.	Seon-Kyeong Jang ...88 more... Scott Vrieze	1.47	0	13	6.5	Nat Hum Behav	2022	Dec 1, 2024 (1 month ago)	
35833142  DOI 	Construction and Application of Polygenic Risk Scores in Autoimmune Diseases.	Chachrit Khunsriraksakul ...4 more... Dajiang J Liu	0.86	0	7	3.5	Front Immunol	2022	Dec 29, 2024 (4 minutes ago)	
35178743  DOI 	Assessing reproducibility of high-throughput experiments in the case of missing data.	Roopali Singh ...1 more... Qunhua Li	0.45	0	3	1.5	Stat Med	2022	Dec 28, 2024 (41 minutes ago)	

Notes



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									

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.