







Core Project R03OD030608

Details

Projects	Name	Award	Publications	Software
1R03OD030608-01	Constructing multi-omics regulatory networks for functional variant annotation	\$335,000.00	3	

Publications

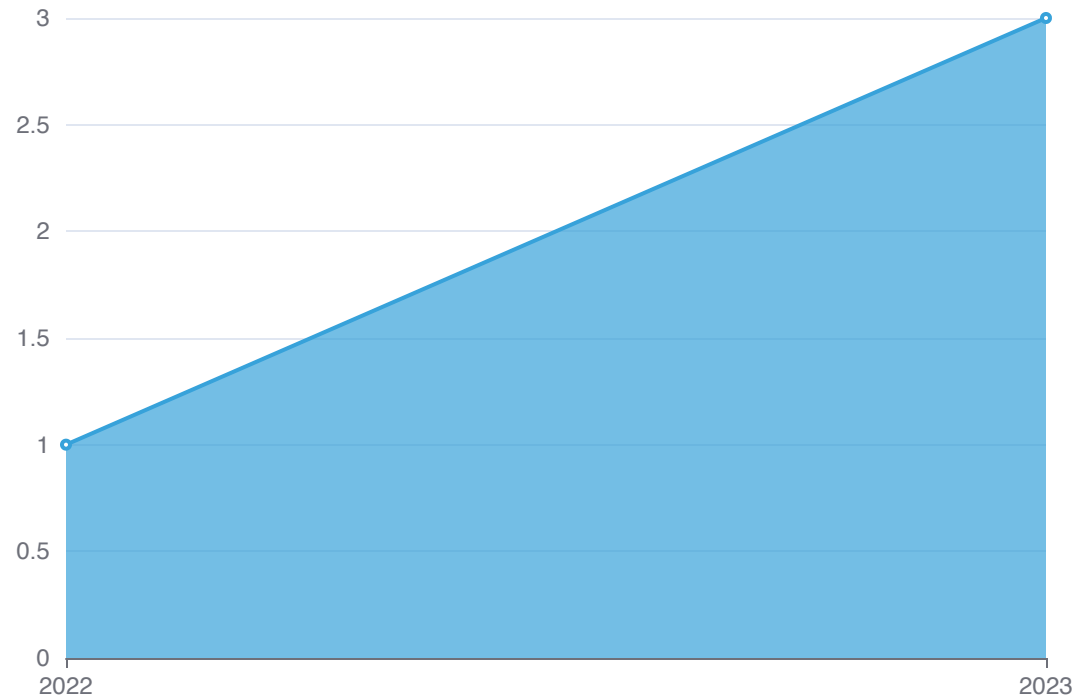
ID	Title	Authors	RCR	SJR	Citations	Cit./year	Journal	Published	Updated
36350676 DOI	FAVOR: functional annotation of variants online resource and annotator for variation across the h...	Hufeng Zhou ...22 more... Xihong Lin	13.73	7.048	36	36	Nucleic Acids Research	2,023	Jun 30, 2024 (4 weeks ago)

36564505  DOI 	Powerful, scalable and resource-efficient meta-analysis of rare variant associations in large who...	Xihao Li ...58 more... Xihong Lin	5.7	17.3	17	17	Nature Genetics	2,023	Jul 12, 2024 (2 weeks ago)
36303018  DOI 	A framework for detecting noncoding rare-variant associations of large-scale whole-genome sequenc...	Zilin Li ...66 more... Xihong Lin	3.95	14.796	34	17	Nature Methods	2,022	Jul 11, 2024 (2 weeks ago)

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

Cumulative Publications



Repositories

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									

PR	Pull (change) request
✓○	Closed/open
Avg Issue/PR	Average time issues/pull requests stay open for before being closed

Notes

Only the `main` (or default) branch is considered (e.g. for # of commits).

of dependencies is totaled from all manifests in repo, direct and transitive, e.g. `package.json` + `package-lock.json`.