

b Core Project R030D030608

Details

Projects	Name	Award	Publications	Repositories
1R03OD030608-01	Constructing multi- omics regulatory networks for functional variant annotation	\$335,000.00	3	0

Publications

Published works associated with this project.

	Title	Authors	RCR	SJR	Citation s	Cit./yea r	Journal	Publishe d	Update
350676 亿 1 亿	FAVOR: functional annotation of variants online resource and annotator for variation across the h	Hufeng Zhou 22 more Xihong Lin	12.4 6	7.048	36	36	Nucleic Acids Research	2,023	Jul 28, 20 (4 week ago)

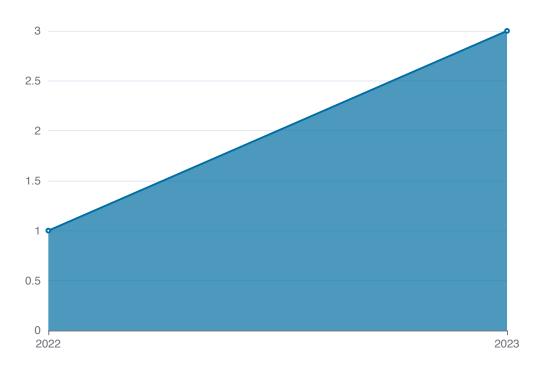
564505 (2 <u>l</u> (2	Powerful, scalable and resource- efficient meta-analysis of rare variant associations in large who	Xihao Li 58 more Xihong Lin	4.68	17.3	16	16	Nature Genetics	2,023	Jul 28, 20 (4 week ago)
<u>803018</u> ♂ <u>L</u> ♂	A framework for detecting noncoding rare-variant associations of large-scale whole-genome sequenc	Zilin Li 66 more Xihong Lin	4.05	14.79 6	36	18	Nature Methods	2,022	Jul 28, 20 (4 week ago)

Notes

RCR = Relative Citation Ratio

SJR = Scimago Journal Rank

Cumulative Publications



Repositories

Software repositories associated with this project.

me	Description	Stars	Watchers	Forks	Issues	PRs	Commits	Contrib.
No data								

Tags Last Commit Avg Issue Avg PR Languages License Readme Contributing Dependencies

No data

Notes

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PR = Pull (change) request

✓ ○ = Closed/open

Avg Issue/PR = Average time issues/pull requests stay open for before being closed

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.

Analytics

Traffic metrics of public websites associated with this project.

Properties

Notes

Active Users = Distinct users who visited the website

New Users = <u>Users who visited the website for the first time</u>

Engaged Sessions = <u>Visits that had significant interaction</u>.

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

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