



(Core Project R03OD030608)

Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD030608-01	Constructing multi-omics regulatory networks for functional variant annotation	\$335K	3 publications	0 repositories	0 properties

Publications

Published works associated with this project.

ID	Title	Author(s)	RCR	SJR	Citations	Cit./year	Journal	Published	Updated
36350676 	FAVOR: functional annotation of variants online resource and annotator for variation across the human genome	Zhou, Hufeng ...23 more... Lin, Xihong	10.073	7.776	106	35.33	Nucleic acids research	2023	Feb 1, 2026
36303018 	A framework for detecting noncoding rare-variant associations of large-scale whole-genome sequencing data	Li, Zilin ...68 more... Lin, Xihong	5.043	17.251	82	20.5	Nature methods	2022	Feb 1, 2026
36564505 	Powerful, scalable and resource-efficient meta-analysis of rare variant associations in large wholegenomes	Li, Xihao ...59 more... Lin, Xihong	3.499	16.586	42	14	Nature genetics	2023	Feb 1, 2026

Publications (cumulative)

Total: 3



Notes

RCR [Relative Citation Ratio ↗](#)

SJR [Scimago Journal Rank ↗](#)

</> Repositories

Software repositories associated with this project.

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Description

	T	S	F	W	C	I	P	R	C	O	C	n	o	L	C	L
	a	Last Commit	a	r	t	s	u	R	t	d	i	o	L	o	n	g
g	g	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15
s	s	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15	2023-01-15

Built on Feb 3, 2026

Developed with support from NIH Award [U54 OD036472](#)

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.

Issue/PR Avg 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

Website metrics 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.