



(Core Project R03OD034494)

Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD034494-01	Using Common Fund datasets for prioritization of disease-associated genetic variants	\$366K	2 publications	0 repositories	0 properties

Publications

Published works associated with this project.

ID	Title	Author s	RC R	SJ R	Citat ions	Cit./ year	Journal	Publi shed	Updat ed
38746164 	Loop Catalog: a comprehensive HiChIP database of human and mouse samples.	Reyna, Joaquin ...8 more... Ay, Ferhat	2.9 99	0	6	6	bioRxiv : the preprint server for biology	2025	Dec 28, 2025
40542429 	Loop Catalog: a comprehensive HiChIP database of human and mouse samples.	Reyna, Joaquin ...8 more... Ay, Ferhat	0	5. 71	3	3	Genome biology	2025	Dec 28, 2025

Publications (cumulative)

Total: 2



Notes

RCR [Relative Citation Ratio ↗](#)

SJR [Scimago Journal Rank ↗](#)

</> Repositories

Software repositories associated with this project.

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Built on Jan 23, 2026

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

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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.