🔬 Core Project R03OD032623

👁 Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD032623-01	Using three-dimensional genome structure to refine eQTL detection	$335K	1 publications	0 repositories	0 properties

📘 Publications

Published works associated with this project.

ID	Title	Authors	RCR	SJR	Citations	Cit./year	Journal	Published	Updated
[38214231](#) ↗ [DOI](#) ↗	Less-is-more: selecting transcription factor binding regions informative for motif inference.	Xu, Jinrui ...2 more... Gerstein, Mark	0.686	7.776	5	2.5	Nucleic acids research	2024	Feb 1, 2026

Notes

RCR	[Relative Citation Ratio](#) ↗
SJR	[Scimago Journal Rank](#) ↗

⟨/⟩ Repositories

Software repositories associated with this project.

Name	Description	Tags	Last Commit	Stars	Forks	Watchers	Commits	Issues	PRs	Issue Avg	PR Avg	Readme	Contributing	Code of Con.	License	Contrib.	Languages

No data

Notes

Repository   For storing, tracking changes to, and collaborating on a piece of software.

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.