



(Core Project R03OD032623)

Overview

High-level info about this project.

| Projects | Name | Award | Publications | Repositories |
|-----------------|---|--------------|----------------|----------------|
| 1R03OD032623-01 | Using three-dimensional genome structure to refine eQTL detection | \$335,000.00 | 1 publications | 0 repositories |

Analytics

0 properties

Publications

Published works associated with this project.

| ID | Title | Author(s) | RC R | SJ R | Citations | Cit./year | Journal | Published | Updated |
|--|---|--|-----------|------|-----------|-----------|-------------------|-----------|------------------------|
| 38214231  | Less-is-more: selecting transcription factor binding regions informative for motif inference. | Xu, Jinrui ...2 more... Gerstein, Mark | 0.7 96 | 0 | 5 | 5 | Nucleic Acids Res | 2024 | Dec 9, 2025 (just now) |

Notes

RCR Relative Citation Ratio 

SJR [Scimago Journal Rank](#) 

</> Repositories

Software repositories associated with this project.

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.