



Core Project R03OD032623



Details

Projects

1R03OD032623-01

Name

Using three-dimensional genome structure to refine eQTL detection

Award

\$335,000.00

Publications

1 publications

Repositories

0 repositories

Analytics

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
38214231 DOI ↗	Less-is-more: selecting transcription factor binding regions informative for motif inference.	Xu, Jinrui ...2 more... Gerstein, Mark	0.796	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.

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

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

Avg Issue/PR 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

Traffic metrics of websites 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.