

L Core Project R030D032623

O Details

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

Publications

Published works associated with this project.

ID	Title	Authors	RC R	SJR	Citatio ns	Cit./ye ar	Journal	Publish ed	Updated
38214231 🗗	Less-is-more: selecting transcription factor binding regions informative for motif inference.	Jinrui Xu 2 more Mark Gerstein	0	7.04 8	0	0	Nucleic Acids Research	2,024	Jul 28, 2024 (4 weeks ago)

Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank

</> Repositories

Software repositories associated with this project.

Name	De	scription	Stars	Watcher	rs Forks	Issue	es 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 2.

New Users <u>Users who visited the website for the first time</u> **.**

Engaged Sessions <u>Visits that had significant interaction</u> **(19)**.

"Top" metrics are measured by number of engaged sessions.