

▶ Core Project R03OD032623

O Details

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

Publications

Published works associated with this project.

ID	Title	Author s	R C R	SJ R	Citat ions	Cit./ year	Journal	Publi shed	Updat ed
38214231 🗹	Less-is-more: selecting transcription factor binding regions informative for motif inference.	Jinrui Xu 2	0	7.0 48	2	2	Nucleic Acids Research	2,024	Oct 26, 2024

more	(1
Mark	week
Gerstei	ago)
n	

Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank

</> Repositories

Software repositories associated with this project.

Name	De	scription	Stars	Watcher	s Forks	Issue	s 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 <u>Distinct users who visited the website</u> 2.

New Users <u>Users who visited the website for the first time</u> 2.

Engaged Sessions <u>Visits that had significant interaction</u> **?**.

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

Generated on Nov 3, 2024

Developed with support from NIH Award U54 OD036472