

# **L** Core Project R03OD032623

#### 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	Author s	R C R	SJ R	Citat ions	Cit./ year	Journal	Publi shed	Updat ed
38214231 <b>乙</b> DOI <b>乙</b>	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	Sep 28, 2024

more	(3
Mark	weeks
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	Ava DD	Languagos	Liconso	Readme	Contributing	Dependencies
Name	rags	Last Commit	Avg issue	Avg PR	Languages	Licerise	Reaume	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> **2**.

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

Generated on Oct 22, 2024

Developed with support from NIH Award U54 OD036472