Core Project R03OD032623

Details

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

Publications

Published works associated with this project.

	Title	Authors	RCR	SJR	Citations	Cit./year	Journal	Published	Update
214231 ⧉ ⧉	Less-is-more: selecting transcription factor binding regions informative for motif inference.	Jinrui Xu …2 more… Mark Gerstein	0	7.048	0	0	Nucleic Acids Research	2,024	Jul 28, 20 (4 week ago)

Notes

RCR = [Relative Citation Ratio ↗](#)

SJR = [Scimago Journal Rank ↗](#)

Repositories

Software repositories associated with this project.

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

me	Tags	Last Commit	Avg Issue	Avg PR	Languages	License	Readme	Contributing	Dependencie
					No data				

Notes

PR   = Pull (change) request

✔ ○   = Closed/open

Avg Issue/PR = Average time issues/pull requests stay open for before being closed

Only the `main` (or default) branch is considered (e.g. for # 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 public websites associated with this project.

Properties

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.