Core Project R03OD032623

Details

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

Publications

Published works associated with this project.

Title	Authors	RCR	SJR	Citations	Cit./year	Journal	Published	Update
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)

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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	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.