

# **L** 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 eOTL 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 🗗 DOI 🗗	Less-is-more: selecting transcription factor binding regions informative for motif inference.	Xu, Jinrui 2	0	0	3	3	Nucleic Acids Res	2024	Aug 19, 2025

more	(just
Gerstei	now)
n, Mark	

#### Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank

### </> Repositories

Software repositories associated with this project.

Name	De	scription	Stars	Watchei	rs Forks	Issue	es PRs	Commits	Contrib.
No data									
Name	Tags	Last Commit	Avg Issue	Avg PR	Languages	License	Readme	Contributing	Dependencies
			8		No data				- op

#### 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> **.** 

New Users Users who visited the website for the first time 2.

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

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

Built on Aug 19, 2025

Developed with support from NIH Award U54 OD036472