



## (Core Project R03OD032623)

### Overview

High-level info about this project.

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

## Publications

Published works associated with this project.

ID	Title	Authors	RC R	SJR	Citations	Cit./year	Journal	Published	Updated
<a href="#">38214231</a> 	<a href="#">DOI</a>  Less-is-more: selecting transcription factor binding regions informative for motif inference.	Xu, Jinrui ...2 more... Gerstein, Mark	0.7 38	7.7 76	5	5	Nucleic acids research	2024	Dec 28, 2025

### Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

# </> Repositories

Software repositories associated with this project.

Name	Description	Last Commit	S	F	W	C	Issues	PRs	I	P	R	n	o	L	C	L	
			t	o	a	o			s	R	e	r	d	i	c	o	a
bioRxiv	A preprint server for biological research articles.	2023-09-21T14:45:00Z															

No data

## Notes

Repository For storing, tracking changes to, and collaborating on a piece of software.

Built on Jan 5, 2026

Developed with support from NIH Award [U54 OD036472](#)

ISSUE LIFETIME: Average time issues/pull requests stay open 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

Website metrics 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.