



## (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	Citat ions	Cit./ year	Journal	Publi shed	Updat ed
<a href="#">38214231</a> 	<a href="#">DOI</a>  Less-is-more: selecting transcription factor binding regions informative for motif inference.	Xu, Jinrui ...2 more... Gerstei n, Mark	0.6 86	7.7 76	5	2.5	Nucleic acids research	2024	Feb 1, 2026

### Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

# </> Repositories

Software repositories associated with this project.

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Description

		S	F	W	C		I	P	R	C	O	C	L	C	L
T	Last Commit	t	o	a	o		s	s	R	t	n	o	L	o	a
a		r	tc	tc	m		u	A	e	d	o	c	i	n	g
g		r	k	h	m		A	v	m	u	C	s	i	r	u
s		s	s	e	it		g	g	m	u	C	s	e	b	g
		s	s	rs	s		v	g	e	t	o	e	b	.	s

No data

## Notes

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

Built on Feb 5, 2026

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

ISSUE LIFETIME: 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

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.