



(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
38214231 	DOI  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.

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Description

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

No data

Notes

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

Built on Jan 8, 2026

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

ISSUES / AVG. 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.