



(Core Project R03OD030599)

Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD030599-01	A database for high-resolution chromatin contact maps and human genetic variants	\$265K	1 publications	0 repositories	0 properties

Publications

Published works associated with this project.

ID	Title	Authors	RCR	SJR	Citations	Cit./year	Journal	Published	Updated
35440119  DOI 	Connecting high-resolution 3D chromatin organization with epigenomics.	Feng, Fan ...3 more.. . Liu, Jie	1.0 98	4.7 61	18	6	Nature communications	2022	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	o	a	
a		r	tc	tc	m		u	A	a	r	e	c	n	t	o	n	
g		r	k	h	m		e	v	a	d	b	f	n	r	e	g	
s		s	s	e	it		A	m	m	u	C	s	i	r	g	e	
		s	s	rs	s		v	g	g	t	o	b	i	n	.	g	

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.

Built on Jan 5, 2026

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

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.