



(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	RC R	SJ R	Citations	Cit./year	Journal	Published	Updated
35440119 	DOI  Connecting high-resolution 3D chromatin organization with epigenomics.	Feng, Fan ...3 more... Liu, Jie	1.0 66	0	17	5.667	Nat Commun	2022	-

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.	2023-11-10T14:45:00Z															

No data

Notes

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

Built on Dec 11, 2025

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.