



# 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	Author s	RC R	SJ R	Citati ons	Cit./y ear	Journal	Publis hed	Upda ted
<a href="#">35440119</a>  <a href="#">DOI</a> 	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	Tags	Last Commit	Stars	Forks	Watchers	Commits	Issues	PRs	Issue Avg	PR Avg	Readme	Contributing	Code of Con.	License	Contrib.	Languages
No data																	

## Notes

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

Issue Avg Average time issues/pull requests stay open for before being closed.

Built on Dec 11, 2025 Developed with support from NIH Award [U54 OD036472](#)

Issue Avg 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.