



# 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
<a href="#">35440119</a>  <a href="#">DOI</a> 	Connecting high-resolution 3D chromatin organization with epigenomics.	Feng, Fan ...3 more.. Liu, Jie	1.067	4.761	18	4.5	Nature communications	2022	Feb 1, 2026

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	Contribs	Languages

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 Feb 15, 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.