



## Core Project R03OD030599



### Overview

High-level info about this project.

#### Projects

1R03OD030599-01

#### Name

A database for high-resolution  
chromatin contact maps and human  
genetic variants

#### Award

\$265,495.00

#### Publications

1 publications

#### Repositories



0 repositories

#### Analytics

0 properties

## Publications

Published works associated with this project.

ID	Title	Autho rs	RC R	SJ R	Citati ons	Cit./y ear	Journal	Publis hed	Update d
<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 Commu n	2022	Dec 10, 2025 (just now)

### 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
No data									

Name	Issue Avg	PR Avg	Readme	Contributing	Code of Con.	License	Contrib.	Languages	Dependencies
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.

Closed/Open    Resolved/unresolved.

Issue/PR 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.