



## Core Project R03OD030599



### Details

#### 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	Updat ed
<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 8, 2025 (just now)

### Notes

RCR

[Relative Citation Ratio](#) 

SJR

[Scimago Journal Rank](#) 

### </> Repositories

Software repositories associated with this project.

Name	Description	Stars	Watchers	Forks	Issues	PRs	Commits	Contrib.
No data								

Name	Tags	Last Commit	Avg Issue	Avg PR	Languages	License	Readme	Contributing	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.
Avg Issue/PR	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

Traffic metrics of websites associated with this project.

### Notes

Active Users	<a href="#">Distinct users who visited the website</a>  .
New Users	<a href="#">Users who visited the website for the first time</a>  .
Engaged Sessions	<a href="#">Visits that had significant interaction</a>  .

"Top" metrics are measured by number of engaged sessions.