



## (Core Project R03OD034500)

### Overview

High-level info about this project.

| Projects        | Name  | Award  | Publications   | Repositories   | Analytics    |
|-----------------|---|--------|----------------|----------------|--------------|
| 1R03OD034500-01 | Allele-specific analysis of human epigenome, transcriptome and high-resolution chromatin organization | \$280K | 0 publications | 0 repositories | 0 properties |

## Publications

Published works associated with this project.

| ID      | Title | Authors | RCR | SJR | Citations | Cit./year | Journal | Published | Updated |
|---------|-------|---------|-----|-----|-----------|-----------|---------|-----------|---------|
| No data |       |         |     |     |           |           |         |           |         |

## Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

# </> Repositories

Software repositories associated with this project.



Built on Jan 19, 2026

Developed with support from NIH Award [U54 OD036472](#)

## 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.