



# Core Project R03OD032623



## Details

| Projects        | Name  | Award        | Publications   | Repositories   | Analytics    |
|-----------------|---|--------------|----------------|----------------|--------------|
| 1R03OD032623-01 | Using three-dimensional genome structure to refine eQTL detection | \$335,000.00 | 1 publications | 0 repositories | 0 properties |



## Publications

Published works associated with this project.

| ID  | Title   | Author<br>s                      | RC<br>R   | SJ<br>R | Citat<br>ions | Cit./<br>year | Journal                 | Publi<br>shed | Updat<br>ed                     |
|---|---|----------------------------------|-----------|---------|---------------|---------------|-------------------------|---------------|---------------------------------|
| <a href="#">38214231</a><br><a href="#">DOI</a> | Less-is-more: selecting transcription factor binding regions informative for motif inference. | Xu,<br>Jinrui<br>...2<br>more... | 0.7<br>96 | 0       | 5             | 5             | Nucleic<br>Acids<br>Res | 2024          | Dec 4,<br>2025<br>(just<br>now) |

|  |                |
|--|----------------|
|  | Gerstein, Mark |
|--|----------------|

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        [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.