



## (Core Project R03OD032623)

### Overview

High-level info about this project.

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

## Publications

Published works associated with this project.

| ID   | Title   | Authors  | RC<br>R   | SJR       | Citat<br>ions | Cit./<br>year | Journal                      | Publi<br>shed | Updat<br>ed        |
|--|---|--|-----------|-----------|---------------|---------------|------------------------------|---------------|--------------------|
| <a href="#">38214231</a>  | <a href="#">DOI</a>  Less-is-more: selecting transcription factor binding regions informative for motif inference. | Xu,<br>Jinrui<br>...2<br>more...<br>Gerstei<br>n, Mark | 0.7<br>38 | 7.7<br>76 | 5             | 5             | Nucleic<br>acids<br>research | 2024          | Dec<br>28,<br>2025 |

### Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

# </> Repositories

Software repositories associated with this project.

| Name    | Description   | Last Commit          | S                    | F                    | W                    | C                    | Issues               | PRs                  | I                    | P                    | R                    | n                    | o                    | L                    | C                    | L                    |                      |
|---------|---|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|
|         |   |                      | t                    | o                    | a                    | o                    |                      |                      | s                    | R                    | e                    | r                    | d                    | i                    | c                    | o                    | a                    |
| bioRxiv | A preprint server for biological research articles. | 2023-01-12T14:45:00Z |

No data

## Notes

Repository For storing, tracking changes to, and collaborating on a piece of software.

Built on Jan 23, 2026

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

ISSUE LIFETIME: Average time issues/pull requests stay open 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.