



# Core Project R03OD034499



## Details









| Projects        | Name  | Award        | Publications   | Repositories   | Analytics    |
|-----------------|---|--------------|----------------|----------------|--------------|
| 1R03OD034499-01 | Deciphering the 3D genome of pediatric brain tumors | \$391,151.00 | 7 publications | 0 repositories | 0 properties |





## Publications

Published works associated with this project.

| ID  | Title  | Authors                | RC<br>R   | SJ<br>R | Citat<br>ions | Cit./<br>year | Journal            | Publi<br>shed | Upda<br>ted                     |
|---|--|------------------------|-----------|---------|---------------|---------------|--------------------|---------------|---------------------------------|
| <a href="#">38796686</a><br><a href="#">DOI</a> | SuPreMo: a computational tool for streamlining in silico perturbation using sequence-based predic... | Gjoni, Ketrin Pollard, | 1.0<br>31 | 0       | 5             | 5             | Bioinfo<br>rmatics | 2024          | Sep 2,<br>2025<br>(just<br>now) |

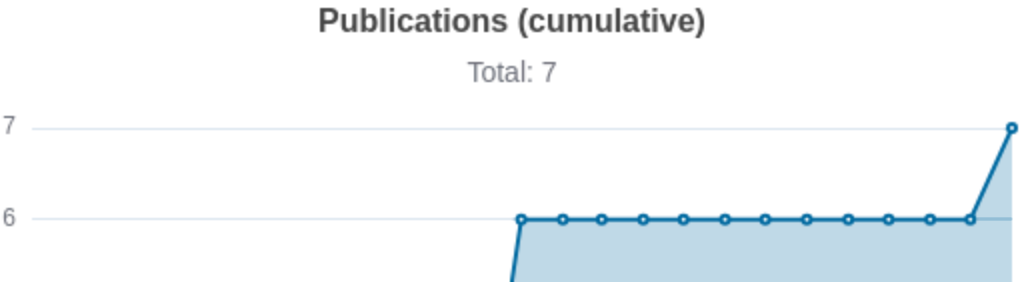
|   |  |   |       |   |   |     |                |      |                        |  |
|---|--|---|-------|---|---|-----|----------------|------|------------------------|--|
|   |  | Katherine S   |       |   |   |     |                |      |                        |  |
| <a href="#">37066196</a> <br><a href="#">DOI</a>      | Comparing chromatin contact maps at scale: methods and insights.                                     | Gunsalus, Laura M<br>...5 more...<br>Pollard, Katherine S | 0.526 | 0 | 6 | 3   | bioRxiv        | 2023 | Sep 2, 2025 (just now) |  |
| <a href="#">37292728</a> <br><a href="#">DOI</a>      | Comparing chromatin contact maps at scale: methods and insights.                                     | Gunsalus, Laura M<br>...5 more...<br>Pollard, Katherine S | 0.229 | 0 | 3 | 1.5 | Res Sq         | 2023 | Sep 2, 2025 (just now) |  |
| <a href="#">39572737</a> <br><a href="#">DOI</a>  | Pooled CRISPR screens with joint single-nucleus chromatin accessibility and transcriptome profiling. | Yan, Rachel E<br>...14 more...<br>Sanjana, Neville E      | 0     | 0 | 4 | 4   | Nat Biotechnol | 2024 | Sep 2, 2025 (just now) |  |
| <a href="#">39574698</a> <br><a href="#">DOI</a>  | De novo structural variants in autism spectrum disorder disrupt distal regulatory interactions of... | Gjoni, Ketrin<br>...3 more...<br>Pollard,                 | 0     | 0 | 1 | 1   | bioRxiv        | 2024 | Sep 2, 2025 (just now) |  |

|  |  |   |   |   |   |   |              |      |                        |  |
|--|--|---|---|---|---|---|--------------|------|------------------------|--|
|  |  | Katherine S                                     |   |   |   |   |              |      |                        |  |
| <a href="#">40108448</a>  | Comparing chromatin contact maps at scale: methods and insights.                                     | Gjoni, Ketrin ...5 more... Pollard, Katherine S | 0 | 0 | 3 | 3 | Nat Metho ds | 2025 | Sep 2, 2025 (just now) |  |
| <a href="#">37961123</a>  | SuPreMo: a computational tool for streamlining <i>in silico</i> perturbation using sequence-based... | Gjoni, Ketrin Pollard, Katherine S              | 0 | 0 | 0 | 0 | bioRxiv      | 2023 | Sep 2, 2025 (just now) |  |

Notes

RCR [Relative Citation Ratio](#)

SJR [Scimago Journal Rank](#)





| 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.

Built on Sep 2, 2025

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