



# 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 | Cita<br>tion<br>s | Cit./<br>year | Journal            | Publi<br>shed | Updat<br>ed        |
|-------------------------------------------------|------------------------------------------------------------------------------------------------------|------------------------|-----------|---------|-------------------|---------------|--------------------|---------------|--------------------|
| <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<br>17,<br>2025 |

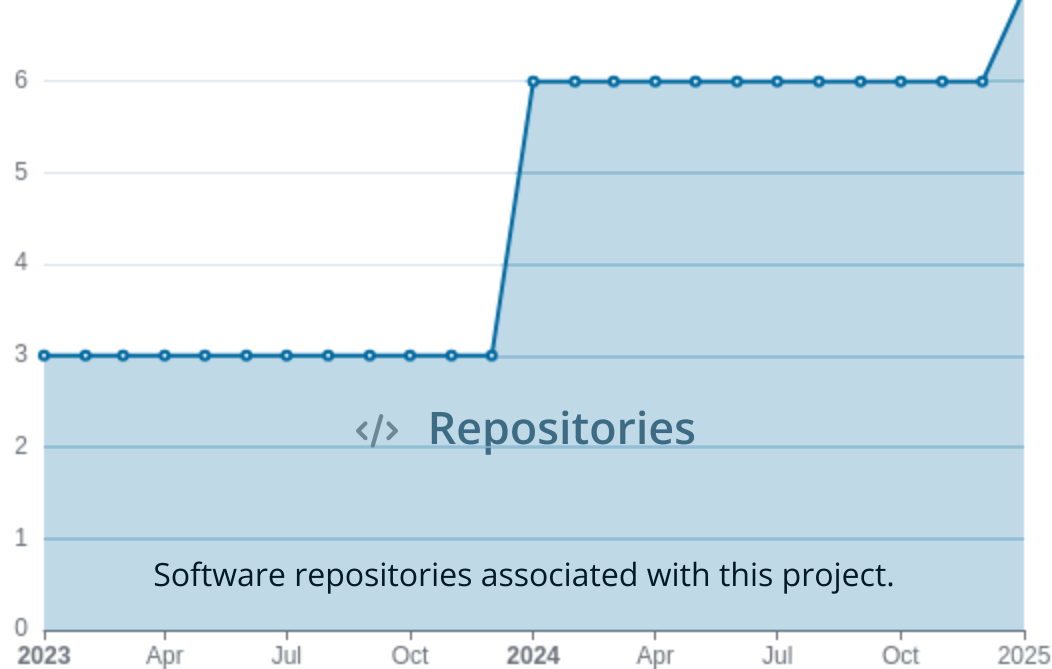
|                                                                                                                                                                                                                         |                                                                                                      |                                                           |       |   |   |     |                |      |                         |            |
|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------|-----------------------------------------------------------|-------|---|---|-----|----------------|------|-------------------------|------------|
|                                                                                                                                                                                                                         |                                                                                                      | Katherine S                                               |       |   |   |     |                |      |                         | (just now) |
| <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 17, 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 17, 2025 (just now) |            |
| <a href="#">37961123</a> <br><a href="#">DOI</a>  | SuPreMo: a computational tool for streamlining <i>in silico</i> perturbation using sequence-based... | Gjoni, Ketrin<br>Pollard, Katherine S                     | 0     | 0 | 0 | 0   | bioRxiv        | 2023 | Sep 17, 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...                            | 0     | 0 | 4 | 4   | Nat Biotechnol | 2024 | Sep 17, 2025 (just now) |            |

|                                                                                                            |                                                                                                      |                                                                   |   |   |   |   |                    |      |                                     |  |
|------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------|---|---|---|---|--------------------|------|-------------------------------------|--|
|                                                                                                            |                                                                                                      | Sanjana,<br>Neville E                                             |   |   |   |   |                    |      |                                     |  |
| <a href="#">40108448</a>  | Comparing chromatin contact maps at scale: methods and insights.                                     | Gjoni,<br>Ketrin<br>...5<br>more...<br>Pollard,<br>Katherine<br>S | 0 | 0 | 3 | 3 | Nat<br>Metho<br>ds | 2025 | Sep<br>17,<br>2025<br>(just<br>now) |  |
| <a href="#">39574698</a>  | De novo structural variants in autism spectrum disorder disrupt distal regulatory interactions of... | Gjoni,<br>Ketrin<br>...3<br>more...<br>Pollard,<br>Katherine<br>S | 0 | 0 | 1 | 1 | bioRxiv            | 2024 | Sep<br>17,<br>2025<br>(just<br>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      [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 17, 2025

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