





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 R | SJ R | Cita tion s | Cit./ year | Journal | Publi shed | Updat ed |
|---|--|------------------------------|-----------|---------|-------------------|---------------|--------------------|---------------|--------------------|
| 38796686  DOI  | SuPreMo: a computational tool for streamlining in silico perturbation using sequence-based predic... | Gjoni, Ketrin Pollard, | 0.9 74 | 0 | 5 | 5 | Bioinfo rmatics | 2024 | Oct 11, 2025 |

| | | | | | | | | | | |
|---|--|---|-------|---|---|-----|---------|------|-------------------------|------------|
| | | Katherine S | | | | | | | | (just now) |
| 37066196  DOI  | Comparing chromatin contact maps at scale: methods and insights. | Gunsalus, Laura M ...5 more... Pollard, Katherine S | 0.502 | 0 | 6 | 3 | bioRxiv | 2023 | Oct 11, 2025 (just now) | |
| 37292728  DOI  | Comparing chromatin contact maps at scale: methods and insights. | Gunsalus, Laura M ...5 more... Pollard, Katherine S | 0.219 | 0 | 3 | 1.5 | Res Sq | 2023 | Oct 11, 2025 (just now) | |
| 37961123  DOI  | 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 | Oct 11, 2025 (just now) | |
| 39574698  DOI  | De novo structural variants in autism spectrum disorder disrupt distal regulatory interactions of... | Gjoni, Ketrin ...3 more... Pollard, | 0 | 0 | 1 | 1 | bioRxiv | 2024 | Oct 11, 2025 (just now) | |

| | | | | | | | | | | |
|--|--|---|---|---|---|---|-----------------|------|-------------------------|--|
| | | Katherine S | | | | | | | | |
| 40108448  | Comparing chromatin contact maps at scale: methods and insights. | Gjoni, Ketrin ...5 more... Pollard, Katherine S | 0 | 0 | 3 | 3 | Nat Metho ds | 2025 | Oct 11, 2025 (just now) | |
| 39572737  | Pooled CRISPR screens with joint single-nucleus chromatin accessibility and transcriptome profiling. | Yan, Rachel E ...14 more... Sanjana, Neville E | 0 | 0 | 4 | 4 | Nat Biotech nol | 2024 | Oct 11, 2025 (just now) | |

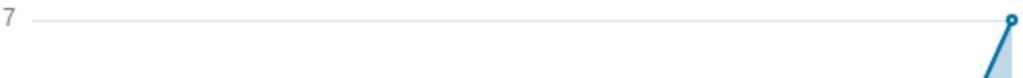
Notes

RCR [Relative Citation Ratio](#) [↗](#)

SJR [Scimago Journal Rank](#) [↗](#)

Publications (cumulative)

Total: 7





| 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 Oct 11, 2025

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