

L Core Project R03OD034499

O 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, | 1.0 31 | 0 | 5 | 5 | Bioinfo rmatics | 2024 | Sep 27, 2025 |

| | | Katherine S | | | | | | | (just now) |
|-----------------------------------|---|---|-----------|---|---|-----|---------|------|-------------------------------------|
| 37066196 🗹 DOI 🗹 | Comparing chromatin contact maps at scale: methods and insights. | Gunsalus, Laura M 5 more Pollard, Katherine S | 0.5 26 | 0 | 6 | 3 | bioRxiv | 2023 | Sep 27, 2025 (just now) |
| 37292728 ② DOI ② | Comparing chromatin contact maps at scale: methods and insights. | Gunsalus, Laura M 5 more Pollard, Katherine S | 0.2 29 | 0 | 3 | 1.5 | Res Sq | 2023 | Sep 27, 2025 (just now) |
| 37961123 🗗 | 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 27, 2025 (just now) |
| 39574698 🗹 | De novo structural variants in autism spectrum disorder disrupt distal regulatory interactions of | Gjoni, Ketrin 3 more Pollard, | 0 | 0 | 1 | 1 | bioRxiv | 2024 | Sep 27, 2025 (just now) |

| | | Katherine S | | | | | | | |
|-----------------------------------|---|---|---|---|---|---|-----------------------|------|-------------------------------------|
| 40108448 乙 DOI 乙 | 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 27, 2025 (just now) |
| 39572737 🗹 DOI 🗹 | 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 | Sep 27, 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 <u>Distinct users who visited the website</u> **.**

New Users <u>Users who visited the website for the first time</u> **.**

Engaged Sessions <u>Visits that had significant interaction</u> **?**.

"Top" metrics are measured by number of engaged sessions.

Built on Sep 27, 2025

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