



(Core Project R03OD038390)

Details

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD038390-01	Ultra-high resolution 3D genome maps for multiple human tissues	\$308,658.00	10 publications	0 repositories	0 properties

Publications

Published works associated with this project.

ID	Title	Authors	RC R	SJ R	Cita tion s	Cit./ yea r	Journal	Publ ishe d	Upda ted
38413840	Enhancer selectivity in space and time: from enhancer-promoter interactions to promoter activation.	Yang, Jin H	14. 82 3	0	69	69	Nat Rev Mol Cell Biol	2024	Oct 15, 2025

		Hansen, Anders S	(just now)
39345388	Dynamics of microcompartment formation at the mitosis-to-G1 transition.	Goel, Viraat Y ...7 more... Hansen, Anders S	Oct 15, 2025 (just now)
40359646	Editorial overview: Emerging perspectives in genome architecture and gene regulation.	Nollmann , Marcelo Hansen, Anders S	Curr Opin Genet Dev 2025 2025 (just now)
40463058	Chromatin Dynamics are Highly Subdiffusive Across Seven Orders of Magnitude.	Mazzocca , Matteo ...3 more... Hansen, Anders S	Oct 15, 2025 (just now)
40081334	Distance matters: How protein regulators facilitate enhancer-promoter interactions and transcript...	Nagano, Masahiro Hansen, Anders S	Oct 15, 2025 (just now)

40654659	Genome structure mapping with high-resolution 3D genomics and deep learning.	Hong, Clarice K Y ...3 more... Hansen, Anders S	0	0	0	0	bioRxiv	2025	Oct 15, 2025 (just now)
39229045	LDB1 establishes multi-enhancer networks to regulate gene expression.	Aboreden, Nicholas G ...11 more... Blobel, Gerd A	0	0	2	2	bioRxiv	2024	Oct 15, 2025 (just now)
39708803	Putative looping factor ZNF143/ZFP143 is an essential transcriptional regulator with no looping f...	Narducci, Domenic N Hansen, Anders S	0	0	12	12	Mol Cell	2025	Oct 15, 2025 (just now)
39935886	Genome-wide absolute quantification of chromatin looping.	Jusuf, James M ...7 more... Hansen, Anders S	0	0	5	5	bioRxiv	2025	Oct 15, 2025 (just now)

[40462903](#) ↗
[DOI](#) ↗

Developing a general AI model for integrating diverse genomic modalities and comprehensive genomi...

Zhang,
Zhenhao
...8
more...
Liu, Jie

0 0 0 0

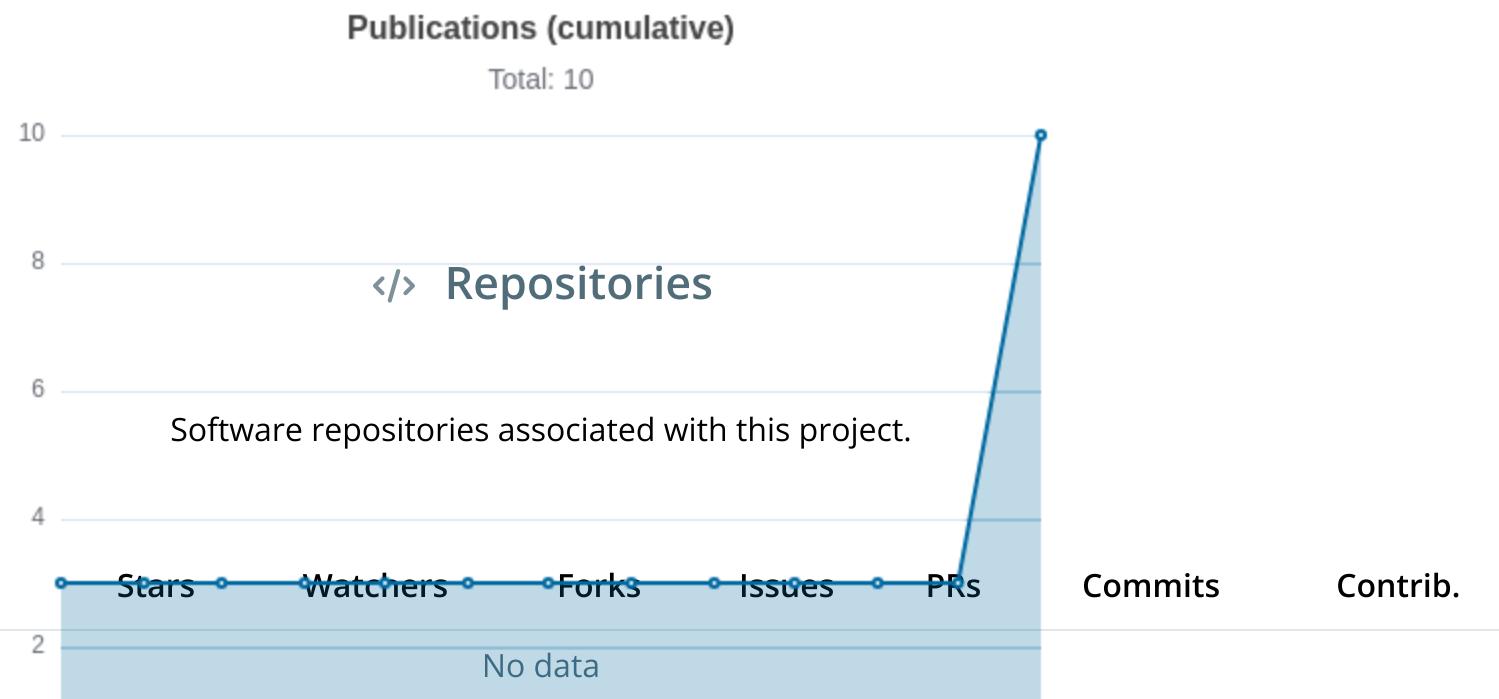
bioRxiv

2025 2025
(just now)

Notes

RCR [Relative Citation Ratio](#) ↗

SJR [Scimago Journal Rank](#) ↗



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 15, 2025

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