



(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	12.	0	79	79	Nat Rev Mol Cell Biol	2024	Dec 5, 2025

	Hansen, Anders S	(just now)
39708803	Putative looping factor ZNF143/ZFP143 is an essential transcriptional regulator with no looping f...	Narducci, Domenic N Hansen, Anders S 6.4 52 0 13 13 Mol Cell 2025 2025 (just now)
39935886	Genome-wide absolute quantification of chromatin looping.	Jusuf, James M ...7 more... Hansen, Anders S 3.4 65 0 7 7 bioRxiv 2025 2025 (just now)
39345388	Dynamics of microcompartment formation at the mitosis-to-G1 transition.	Goel, Viraat Y ...7 more... Hansen, Anders S 0.5 5 0 4 4 bioRxiv 2024 2025 (just now)
39229045	LDB1 establishes multi-enhancer networks to regulate gene expression.	Aboreden , Nicholas G ...11 more... Blobel, Gerd A 0.4 36 0 3 3 bioRxiv 2024 2025 (just now)

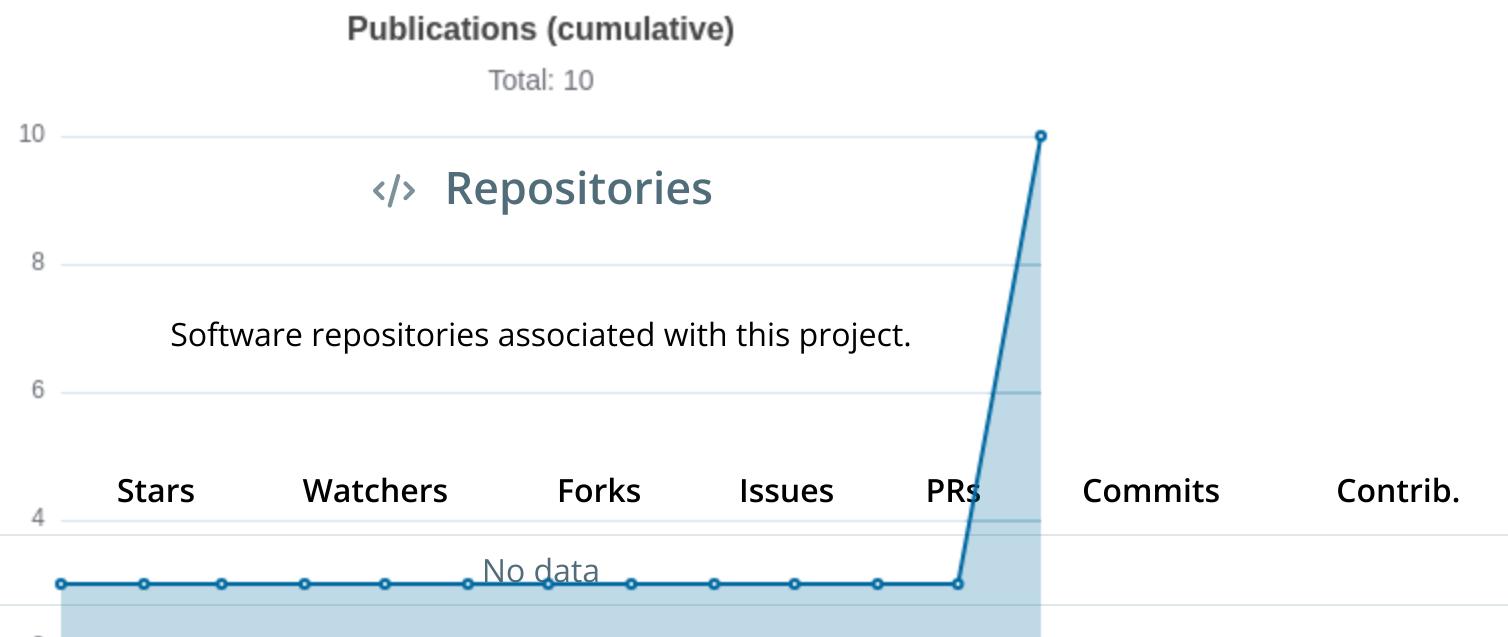
40462903	Developing a general AI model for integrating diverse genomic modalities and comprehensive genomi...	Zhang, Zhenhao ...8 more... Liu, Jie	0	0	0	0	bioRxiv	2025	Dec 5, 2025 (just now)
40463058	Chromatin Dynamics are Highly Subdiffusive Across Seven Orders of Magnitude.	Mazzocca, Matteo ...3 more... Hansen, Anders S	0	0	2	2	bioRxiv	2025	Dec 5, 2025 (just now)
40359646	Editorial overview: Emerging perspectives in genome architecture and gene regulation.	Nollmann , Marcelo Hansen, Anders S	0	0	0	0	Curr Opin Genet Dev	2025	Dec 5, 2025 (just now)
40081334	Distance matters: How protein regulators facilitate enhancer-promoter interactions and transcript...	Nagano, Masahiro Hansen, Anders S	0	0	1	1	Cell Genom	2025	Dec 5, 2025 (just now)

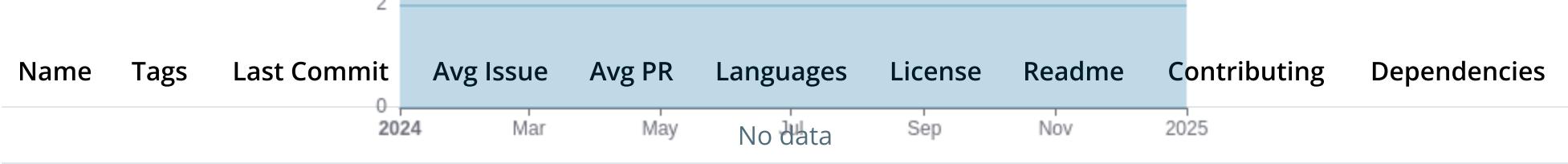
40654659	Genome structure mapping with high-resolution 3D genomics and deep learning.	Hong, Clarice K Y ...3 more... Hansen, Anders S	0	0	1	1	bioRxiv	2025	Dec 5, 2025 (just now)
--------------------------	--	---	---	---	---	---	---------	------	------------------------

Notes

RCR [Relative Citation Ratio](#)

SJR [Scimago Journal Rank](#)





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 Dec 5, 2025

Developed with support from NIH Award [U54 OD036472](#)