

# **L** Core Project R03OD038390

### O Details

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

## 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 🗗 DOI 🗗	Enhancer selectivity in space and time: from enhancer-promoter interactions to promoter activation.	Yang, Jin H	14. 63 5	0	64	64	Nat Rev Mol Cell Biol	2024	Sep 20, 2025

		Hansen, Anders S							(just now)
40081334 🖸	Distance matters: How protein regulators facilitate enhancer-promoter interactions and transcript	Nagano, Masahiro Hansen, Anders S	0	0	0	0	Cell Genom	2025	Sep 20, 2025 (just now)
39935886 ☑ DOI ☑	Genome-wide absolute quantification of chromatin looping.	Jusuf, James M7 more Hansen, Anders S	0	0	3	3	bioRxiv	2025	Sep 20, 2025 (just now)
40463058 <b>乙</b> DOI <b>乙</b>	Chromatin Dynamics are Highly Subdiffusive Across Seven Orders of Magnitude.	Mazzocca , Matteo 3 more Hansen, Anders S	0	0	0	0	bioRxiv	2025	Sep 20, 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	Sep 20, 2025 (just now)

39229045 🗗 DOI 🗗	LDB1 establishes multi-enhancer networks to regulate gene expression.	Aboreden , Nicholas G 11 more Blobel, Gerd A	0	0	2	2	bioRxiv	2024	Sep 20, 2025 (just now)
39345388 🗗 DOI 🗹	Dynamics of microcompartment formation at the mitosis-to-G1 transition.	Goel, Viraat Y 7 more Hansen, Anders S	0	0	3	3	bioRxiv	2024	Sep 20, 2025 (just now)
39708803 🗗 DOI 🗗	Putative looping factor ZNF143/ZFP143 is an essential transcriptional regulator with no looping f	Narducci, Domenic N Hansen, Anders S	0	0	11	11	Mol Cell	2025	Sep 20, 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	Sep 20, 2025 (just now)
40654659 ♂ DOI ♂	Genome structure mapping with high- resolution 3D genomics and deep learning.	Hong, Clarice K Y	0	0	0	0	bioRxiv	2025	Sep 20, 2025



### Notes

Name

RCR Relative Citation Ratio

SJR Scimago Journal Rank

Description

### Publications (cumulative)



Tul

May

Sen

Nov

Contrib.

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 2.

New Users Users who visited the website for the first time 2.

Engaged Sessions Visits that had significant interaction <a>\mathbb{Z}</a>.

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

Built on Sep 20, 2025

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