



## (Core Project R03OD038390)

### Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD038390-01	Ultra-high resolution 3D genome maps for multiple human tissues	\$309K	12 publications	0 repositories	0 properties

## Publications

Published works associated with this project.

ID	Title	Authors	RC R	SJR	Cita tion s	Cit./ yea r	Journal	Publ ishe d	Upda ted
<a href="#">38413840</a> 	Enhancer selectivity in space and time: from enhancer-promoter interactions to promoter activation.	Yang, Jin H Hansen, Anders S	13. 16 6	37. 35 3	91	91	Nature reviews. Molecular cell biology	2024	Dec 28, 2025
<a href="#">39708803</a> 	Putative looping factor ZNF143/ZFP143 is an essential transcriptional regulator with no looping f...	Narducci, Domenic N Hansen, Anders S	6.4 98	9.0 51	13	13	Molecular cell	2025	Dec 28, 2025
<a href="#">39935886</a> 	Genome-wide absolute quantification of chromatin looping.	Jusuf, James M ...7 more... Hansen, Anders S	4.4 98	0	9	9	bioRxiv : the preprint server for biology	2025	Dec 28, 2025
<a href="#">39345388</a> 	Dynamics of microcompartment formation at the mitosis-to-G1 transition.	Goel, Viraat Y ...7 more...	0.5 1	0	4	4	bioRxiv : the preprint server for biology	2024	Dec 28, 2025

	Hansen, Anders S							
<a href="#">39229045</a>	LDB1 establishes multi-enhancer networks to regulate gene expression.	Aboreden , Nicholas G...11 more... Blobel, Gerd A	0.4 05	0	3	3	bioRxiv : the preprint server for biology	2024 Jan 23, 2026
<a href="#">40654659</a>	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 : the preprint server for biology	2025 Dec 28, 2025
<a href="#">41277691</a>	Developing a general AI model for integrating diverse genomic modalities and comprehensive genomi...	Zhang, Zhenhao ...11 more... Liu, Jie	0	7.7 76	0	0	Nucleic acids research	2025 Dec 28, 2025
<a href="#">40462903</a>	Developing a general AI model for integrating diverse genomic modalities and comprehensive genomi...	Zhang, Zhenhao ...8 more... Liu, Jie	0	0	0	0	bioRxiv : the preprint server for biology	2025 Dec 28, 2025

[41107486](#) [DOI](#)

Dynamics of microcompartment formation at the mitosis-to-G1 transition.

Goel,

Viraat Y

...7

more...

Hansen,  
Anders S

0

0

3

3

Nat Struct Mol  
Biol

2025

Dec  
28,  
2025

[40081334](#) [DOI](#)

Distance matters: How protein regulators facilitate enhancer-promoter interactions and transcript...

Nagano,  
Masahiro  
Hansen,  
Anders S

0

6.2  
38

1

1

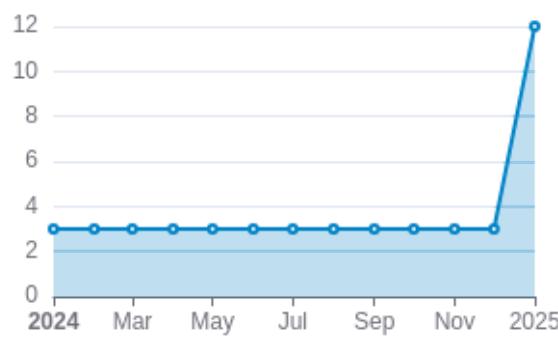
Cell genomics

2025

Dec  
28,  
2025

### Publications (cumulative)

Total: 12



### Notes

RCR [Relative Citation Ratio](#)

SJR [Scimago Journal Rank](#)

## </> Repositories

Software repositories associated with this project.

Name	Description	Last Commit	S	F	W	C	I	P	R	C	O	C	N	O	L	C	L
			t	o	a	m	s	e	A	R	e	n	o	d	i	c	o
			at	to	tc	m	re	it	Ad	re	ri	on	o	de	ci	on	g

No data

## Notes

Repository     For storing, tracking changes to, and collaborating on a piece of software.

Built on Jan 25, 2026

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

ISSUE COUNT - 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

Website metrics 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.