



(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.

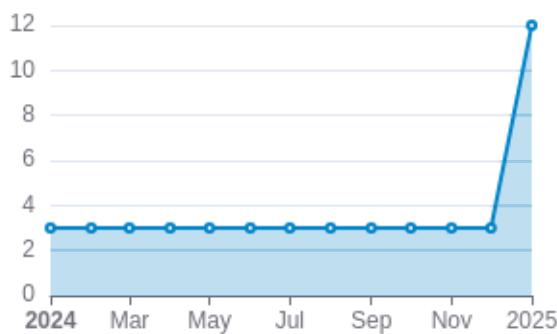
ID	Title	Authors	RC R	SJR	Cita tion s	Cit./ yea r	Journal	Publ ishe d	Upda ted
38413840	Enhancer selectivity in space and time: from enhancer-promoter interactions to promoter activation.	Yang, Jin H Hansen, Anders S	13. 16 6	37. 35 3	91	91	Nature reviews. Molecular cell biology	2024	Dec 28, 2025
39708803	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
39935886	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
39345388	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									
39229045	LDB1 establishes multi-enhancer networks to regulate gene expression.	Aborede n, Nicholas G ...11 more...	0.4 05	0	3	3	bioRxiv : the preprint server for biology	2024	Jan 23, 2026	
40654659	Genome structure mapping with high-resolution 3D genomics and deep learning.	Blobel, Gerd A	Hong, Clarice K Y ...3 more...	0	0	1	1	bioRxiv : the preprint server for biology	2025	Dec 28, 2025
41107486	Dynamics of microcompartment formation at the mitosis-to-G1 transition.	Hansen, Anders S	Goel, Viraat Y ...7 more...	0	0	3	3	Nat Struct Mol Biol	2025	Dec 28, 2025

40359646	Editorial overview: Emerging perspectives in genome architecture and gene regulation.	Nollman n, Marcelo Hansen, Anders S	0	2.0 94	0	0	Current opinion in genetics & development	2025	Dec 28, 2025
41277691	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
40081334	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	Tags	Last Commit	S	F	W	C	I	S	P	R	T	O	C	N	L	C	A
John Doe	Full Stack Developer	js, react, node	2023-10-01 15:30:00 UTC	John	Doe	React	Node.js	Issue	Sub	PR	Review	Test	Open	Completed	Blocked	In Progress	On Hold	Not Started
Jane Smith	Frontend Developer	js, html, css	2023-10-02 10:00:00 UTC	Jane	Smith	HTML/CSS	React	Issue	Sub	PR	Review	Test	Open	Completed	Blocked	In Progress	On Hold	Not Started
Bob Johnson	Backend Developer	node, express, mongo	2023-10-03 12:00:00 UTC	Bob	Johnson	Node.js	Express.js	Issue	Sub	PR	Review	Test	Open	Completed	Blocked	In Progress	On Hold	Not Started
Sarah Williams	Full Stack Developer	js, react, mongo	2023-10-04 09:00:00 UTC	Sarah	Williams	React	MongoDB	Issue	Sub	PR	Review	Test	Open	Completed	Blocked	In Progress	On Hold	Not Started

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.

Built on Jan 28, 2026

Developed with support from NIH Award U54 OD036472

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.