



## (Core Project R03OD036499)

### Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD036499-01	Cell type harmonization of single cell data in HuBMAP and GTEx	\$390K	6 publications	0 repositories	0 properties

## Publications

Published works associated with this project.

ID	Title	Authors	RCR	SJR	Citations	Cit./year	Journal	Published	Updated
<a href="#">40082611</a>	Human BioMolecular Atlas Program (HuBMAP): 3D Human Reference Atlas construction and usage.	Börner, Katy ...19 more... Herr, Bruce W	11. 24 6	17. 25 1	23	23	Nature methods	2025	Feb 1, 2026
<a href="#">40893796</a>	Discovery of optimal cell type classification marker genes from single cell RNA sequencing data.	Liu, Angela ...5 more... Zhang, Yun	1.1 3	0	8	4	BMC methods	2024	Feb 1, 2026
<a href="#">38826261</a>	Human BioMolecular Atlas Program (HuBMAP): 3D Human Reference Atlas Construction and Usage.	Börner, Katy ...19 more... Herr, Bruce W	0.9 29	0	5	2.5	bioRxiv : the preprint server for biology	2024	Feb 1, 2026
<a href="#">38712147</a>	Discovery of optimal cell type classification marker genes from single cell RNA	Liu, Angela	0.1 5	0	1	0.5	bioRxiv : the preprint server	2024	Feb 1,

	sequencing data.		...5 more... Zhang, Yun		for biology		2026
<a href="#">40475598</a>	A Multimodal Spatial and Epigenomic Atlas of Human Adult Lung Topography.	Duong, Thu Elizabeth ...21 more... Pryhuber , Gloria S	0 0 1 1	bioRxiv : the preprint server for biology	2025	Feb 1, 2026	
<a href="#">40568082</a>	Benchmarking single cell transcriptome matching methods for incremental growth of reference atlases.	Hu, Joyce ...9 more... Zhang, Yun	0 0 1 1	bioRxiv : the preprint server for biology	2025	Feb 1, 2026	

### Publications (cumulative)

Total: 6



## Notes

RCR [Relative Citation Ratio ↗](#)

SJR [Scimago Journal Rank ↗](#)

# </> Repositories

Built on Feb 3, 2026

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

Name	Description	Last Commit	S	F	W	C	Issues	PRs	I	P	R	L	C	O	C	n	o	L	C	a	
			t	o	a	o			s	R	e	d	i	c	o	n	g	t	o	n	g
bioRxiv	bioRxiv is a preprint server for life sciences research. It is a community-driven platform that allows researchers to share their work before it has been peer-reviewed. The repository contains the source code for the bioRxiv website and various tools used in the preprint submission process.	2023-02-01T12:00:00Z	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100

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.

Issue/PR Avg 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.