



(Core Project R03OD034501)

Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories
1R03OD034501-01	Integration of GTEx and HuBMAP data to gain population-level cell-type-specific insights	\$314,739.00	6 publications	0 repositories

Analytics

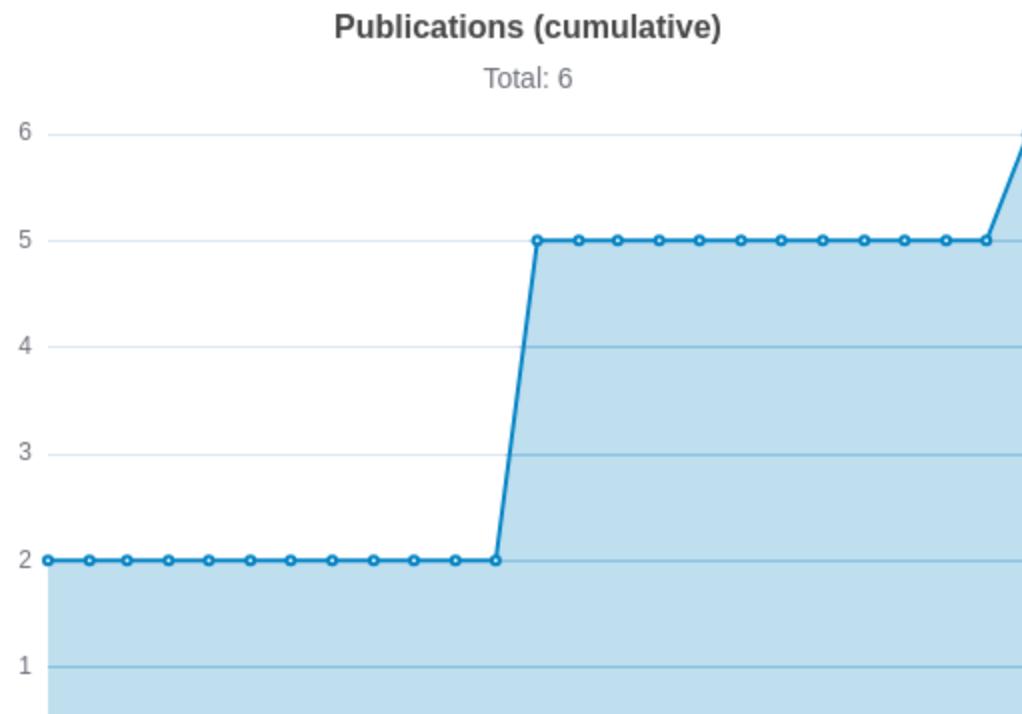
0 properties

Publications

Published works associated with this project.

ID	Title	Authors	RC R	SJ R	Citat ions	Cit./ year	Journal	Publi shed	Upda ted
38168620	scMD facilitates cell type deconvolution using single-cell DNA methylation references.	Cai, Manqi ...2 more... Wang, Jiebiao	9.7 87	0	42	42	Commun Biol	2024	Dec 9, 2025 (just now)
37563770	Transcriptional risk scores in Alzheimer's disease: From pathology to cognition.	Pyun, Jung- Min ...7 more... Nho, Kwangsi k	1.6 17	0	8	8	Alzheimer s Dement	2024	Dec 9, 2025 (just now)
36993280	Accurate estimation of rare cell type fractions from tissue omics data via hierarchical deconvolu...	Huang, Penghui ...3 more... Wang, Jiebiao	0.3 09	0	3	1.5	bioRxiv	2023	Dec 9, 2025 (just now)
37577715	scMD: cell type deconvolution using single-cell DNA methylation references.	Cai, Manqi ...2 more... Wang, Jiebiao	0.2 08	0	2	1	bioRxiv	2023	Dec 9, 2025 (just now)

39149243	BLEND: Probabilistic Cellular Deconvolution with Automated Reference Selection.	Huang, Penghui ...2 more...	0	0	0	0	bioRxiv	2024	Dec 9, 2025 (just now)
41112940	EMixed: Probabilistic Multi-Omics Cellular Deconvolution of Bulk Omics Data.	Cai, Manqi ...5 more...	0	0	0	0	J Data Sci	2025	Dec 9, 2025 (just now)





Notes

RCR Relative Citation Ratio 

SJR [Scimago Journal Rank](#) 

</> Repositories

Software repositories associated with this project.

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.

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