

# **▶** Core Project R03OD034501

## Operation

Projects	Name	Award	Publications	Repositories	Analytics
1R03OD034501-01	Integration of GTEx and HuBMAP data to gain population-level cell-type-specific insights	\$314,739.00	5 publications	0 repositories	0 properties

## Publications

Published works associated with this project.

ID	Title	Author s	R C R	SJ R	Cita tion s	Cit./ yea r	Journal	Publi shed	Updat ed
36993280 <b>♂</b> DOI <b>♂</b>	Accurate estimation of rare cell type fractions from tissue omics data via	Penghu i Huang	0	0	1	1	bioRxiv	2,02 3	Sep 1, 2024

	hierarchical deconvolu	3 more Jiebiao Wang							(3 weeks ago)
38168620 <b>乙</b>	scMD facilitates cell type deconvolution using single-cell DNA methylation references.	Manqi Cai 2 more Jiebiao Wang	0	2.0 9	1	1	Communica tions Biology	2,02 4	Sep 1, 2024 (3 weeks ago)
37577715 🗗 DOI 🗗	scMD: cell type deconvolution using single- cell DNA methylation references.	Manqi Cai 2 more Jiebiao Wang	0	0	0	0	bioRxiv	2,02 3	Sep 1, 2024 (3 weeks ago)
39149243 🖸 DOI 🗗	BLEND: Probabilistic Cellular Deconvolution with Automated Reference Selection.	Penghu i Huang 2 more Jiebiao Wang	0	0	0	0	bioRxiv	2,02 4	Sep 1, 2024 (3 weeks ago)
37563770 CZ	Transcriptional risk scores in Alzheimer's disease: From pathology to cognition.	Jung- Min Pyun 7	0	3.2 26	1	1	Alzheimer's and Dementia	2,02 4	Sep 1, 2024 (3

more	weeks
Kwangs	ago)
ik Nho	

#### Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank

## </> Repositories



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.

Active Users <u>Distinct users who visited the website</u> **.** 

New Users <u>Users who visited the website for the first time</u> **.** 

Engaged Sessions <u>Visits that had significant interaction</u> **?**.

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

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