



Core Project R03OD034501









Details

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
37577715 DOI	scMD: cell type deconvolution using single-cell DNA methylation references.	Manqi Cai	0	0	0	0	bioRxiv	2,023	Sep 1, 2024

		...2 more... Jiebiao Wang								(3 weeks ago)
36993280  DOI 	Accurate estimation of rare cell type fractions from tissue omics data via hierarchical deconvolu...	Penghu i Huang ...3 more... Jiebiao Wang	0	0	1	1	bioRxiv	2,023	Sep 1, 2024 (3 weeks ago)	
37563770  DOI 	Transcriptional risk scores in Alzheimer's disease: From pathology to cognition.	Jung- Min Pyun ...7 more... Kwangs ik Nho	0	3.2 26	1	1	Alzheimer's and Dementia	2,024	Sep 1, 2024 (3 weeks ago)	
38168620  DOI 	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	2,024	Sep 1, 2024 (3 weeks ago)	
39149243  DOI 	BLEND: Probabilistic Cellular Deconvolution with Automated Reference Selection.	Penghu i Huang ...2	0	0	0	0	bioRxiv	2,024	Sep 1, 2024 (3	

more...
Jiebiao
Wang

weeks
ago)

Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

</> Repositories

Software repositories associated with this project.

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.

Generated on Sep 22, 2024

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