

# **L** Core Project R03OD034501

### **Details**

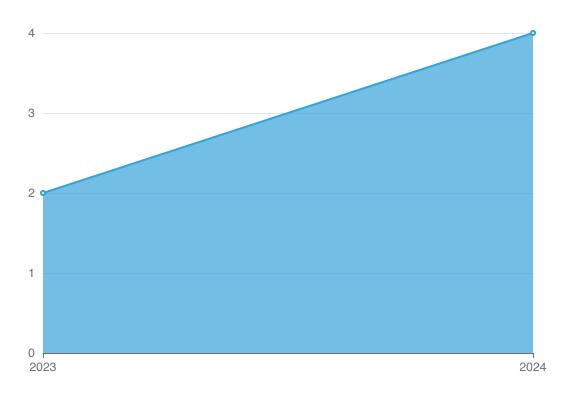
Projects	Name	Award	Publications	Software
1R03OD034501-01	Integration of GTEx and HuBMAP data to gain population-level cell-type-specific insights	\$314,739.00	4	0 repositories

### **Publications**

ID	Title	Author s	R C R	SJ R	Cita tion s	Cit./ year	Journal	Publi shed	Upda ted
36993280 🖸 DOI 🗹	Accurate estimation of rare cell type fractions from tissue omics data via hierarchical deconvolu	Penghu i Huang 3 more	0	0.3 18	1	1	Gene Expression Patterns	2023	Jun 30, 2024

		Jiebiao Wang							
37563770 🗗 DOI 🗗	Transcriptional risk scores in Alzheimer's disease: From pathology to cognition.	Jung- Min Pyun 7 more Kwangs ik Nho	0	2.5 74	1	1	Bioinformati cs	2024	Jun 30, 2024
<u>38168620</u>	scMD facilitates cell type deconvolution using single-cell DNA methylation references.	Manqi Cai 2 more Jiebiao Wang	0	0.9 03	1	1	Metabolites	2024	Jun 30, 2024
37577715 🗗 DOI 🗗	scMD: cell type deconvolution using single- cell DNA methylation references.	Manqi Cai 2 more Jiebiao Wang	0	0.3 18	0	0	Gene Expression Patterns	2023	Jun 30, 2024

#### **Cumulative Publications**



## Repositories

Nam e	Descript ion	_	Sta rs	_	Watch ers	Open Issues	Updat ed	•		Read me	Contribut ing	Dependen cies
No data												