

# **L** Core Project OT2OD036435

#### O Details

Projects	Name	Award	Publications	Repositories	Analytics
1OT2OD036435-01	The CFDE Workbench	\$3,250,000.00	12 publications	0 repositories	0 properties
30T20D036435-01S1					

### Publications

Published works associated with this project.

ID	Title	Authors	RC R	SJ R	Cita tion s	Cit./ yea r	Journal	Publ ishe d	Upda ted
38643247 🗹 DOI 🗗	Rummagene: massive mining of gene sets from supporting materials of biomedical research publicati	Clarke, Daniel J B 4 more	5.0 47	0	16	16	Commun Biol	2024	Aug 24, 2025

		Ma'ayan, Avi							(just now)
39569206 🗗	RummaGEO: Automatic mining of human and mouse gene sets from GEO.	Marino, Giacomo B 3 more Ma'ayan, Avi	2.7 6	0	8	8	Patterns (N Y)	2024	Aug 24, 2025 (just now)
37460679 ♂ DOI ♂	Toxicology knowledge graph for structural birth defects.	Evangelista , John Erol 16 more Ma'ayan, Avi	2.1 22	0	13	6.5	Commun Med (Lond)	2023	Aug 24, 2025 (just now)
37953774 🗷 DOI 🗗	Dex-Benchmark: datasets and code to evaluate algorithms for transcriptomics data analysis.	Xie, Zhuorui 1 more Ma'ayan, Avi	0.1 88	0	1	0.5	PeerJ	2023	Aug 24, 2025 (just now)
38339304 ☑ DOI ☑	Molecular Atlas of HER2+ Breast Cancer Cells Treated with Endogenous Ligands: Temporal Insights i	Mukund, Kavitha 10 more Subramani am, Shankar	0	0	1	1	Cancers (Basel)	2024	Aug 24, 2025 (just now)

39565691 🗹 DOI 🗹	Protocol for using Multiomics2Targets to identify targets and driver kinases for cancer cohorts p	Marino, Giacomo B 6 more Ma'ayan, Avi	0	0	0	0	STAR Protoc	2024	Aug 24, 2025 (just now)
40501705 🗗	A Gene Set Foundation Model Pre-Trained on a Massive Collection of Diverse Gene Sets.	Clarke, Daniel J B 1 more Ma'ayan, Avi	0	0	0	0	bioRxiv	2025	Aug 24, 2025 (just now)
39990419 🖸 DOI 🖸	Spatially distinct cellular and molecular landscapes define prognosis in triple negative breast c	Mukund, Kavitha 13 more Subramani am, Shankar	0	0	0	0	bioRxiv	2025	Aug 24, 2025 (just now)
40308216 🗗	L2S2: chemical perturbation and CRISPR KO LINCS L1000 signature search engine.	Marino, Giacomo B 2 more Ma'ayan, Avi	0	0	1	1	Nucleic Acids Res	2025	Aug 24, 2025 (just now)
40133794 🖸	IncRNAlyzr: Enrichment Analysis for IncRNA Sets.	Evangelista , John Erol 5 more	0	0	2	2	J Mol Biol	2025	Aug 24, 2025

		Ma'ayan, Avi							(just now)
39415993 <b>乙</b> DOI <b>乙</b>	The molecular chronology of mammary epithelial cell fate switching.	Vallmajo- Martin, Queralt 7 more Lytle, Nikki K	0	0	0	0	bioRxiv	2024	Aug 24, 2025 (just now)
38645198 <b>♂</b> DOI <b>♂</b>	RummaGEO: Automatic Mining of Human and Mouse Gene Sets from GEO.	Marino, Giacomo B 2 more Ma'ayan, Avi	0	0	1	1	bioRxiv	2024	Aug 24, 2025 (just now)

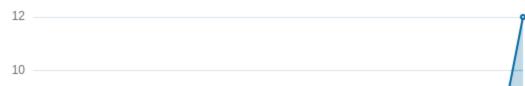
### Notes

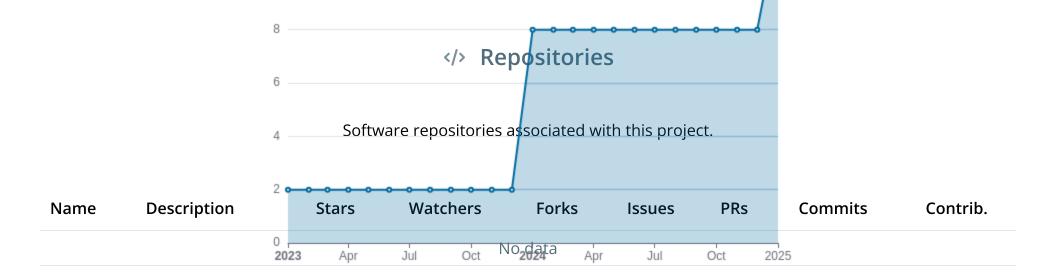
RCR Relative Citation Ratio

SJR Scimago Journal Rank

### Publications (cumulative)

Total: 12





Name	Tags	Last Commit	Avg Issue	Avg PR	Languages	License	Readme	Contributing	Dependencies
					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.

Avg Issue/PR 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

Traffic metrics of websites associated with this project.

#### Notes

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> **?**.

Built on Aug 24, 2025

Developed with support from NIH Award U54 OD036472

<sup>&</sup>quot;Top" metrics are measured by number of engaged sessions.