

# **L** Core Project OT2OD036435

## O Details

Projects	Name	Award	Publications	Repositories	Analytics
1OT2OD036435-01	The CFDE Workbench	\$7,184,710.00	15 publications	0 repositories	0 properties
3OT2OD036435-01S2					
3OT2OD036435-01S1					

# 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 🗹	Rummagene: massive mining of gene sets from supporting materials of biomedical	Clarke, Daniel J B	4.9 09	0	18	18	Commun Biol	2024	Oct 7,

	research publicati	4 more Ma'ayan, Avi							2025 (just now)
39569206 🗗	RummaGEO: Automatic mining of human and mouse gene sets from GEO.	Marino, Giacomo B 3 more Ma'ayan, Avi	2.6 18	0	9	9	Patterns (N Y)	2024	Oct 7, 2025 (just now)
37460679 <b>乙</b>	Toxicology knowledge graph for structural birth defects.	Evangelista , John Erol 16 more Ma'ayan, Avi	2.4 32	0	17	8.5	Commun Med (Lond)	2023	Oct 7, 2025 (just now)
37953774 🗗	Dex-Benchmark: datasets and code to evaluate algorithms for transcriptomics data analysis.	Xie, Zhuorui 1 more Ma'ayan, Avi	0.1 71	0	1	0.5	PeerJ	2023	Oct 7, 2025 (just now)
38645198 <b>乙</b> DOI <b>乙</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	Oct 7, 2025 (just now)

40956607 🗗	Creating an Interactive Web Interface for Networks Stored in Knowledge Graph Databases.	Evangelista , John Erol 5 more Ma'ayan, Avi	0	0	1	1	Curr Protoc	2025	Oct 7, 2025 (just now)
40308216 ☑ DOI ☑	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	Oct 7, 2025 (just now)
40133794 🗗	IncRNAlyzr: Enrichment Analysis for IncRNA Sets.	Evangelista , John Erol 5 more Ma'ayan, Avi	0	0	3	3	J Mol Biol	2025	Oct 7, 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	Oct 7, 2025 (just now)
39990419 🖸	Spatially distinct cellular and molecular landscapes define prognosis in triple negative breast c	Mukund, Kavitha 13 more Subramani	0	0	1	1	bioRxiv	2025	Oct 7, 2025 (just now)

		am, Shankar							
40672200 ☑ DOI ☑	A novel subtyping method for TNBC with implications for prognosis and therapy.	Mesrizade h, Zahra 9 more Subramani am, Shankar	0	0	0	0	bioRxiv	2025	Oct 7, 2025 (just now)
39415993 ☑ DOI ☑	The molecular chronology of mammary epithelial cell fate switching.	Vallmajo- Martin, Queralt 7 more Lytle, Nikki K	0	0	0	0	bioRxiv	2024	Oct 7, 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	Oct 7, 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	Oct 7, 2025 (just now)

40832173 ☑ DOI ☑	ChEA-KG: Human Transcription Factor Regulatory Network with a Knowledge Graph Interactive User In	Byrd, Anna l 4 more Ma'ayan, Avi	0	0	0	0	bioRxiv	2025	Oct 7, 2025 (just now)
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## Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank



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.

# Analytics

Traffic metrics of websites associated with this project.

#### Notes

Active Users <u>Distinct users who visited the website</u> 2.

New Users <u>Users who visited the website for the first time</u> **2**.

Engaged Sessions <u>Visits that had significant interaction</u> **2**.

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