



## (Core Project OT2OD036435)

### Overview

High-level info about this project.

Projects	Name	Award	Publications	Repositories	Analytics
1OT2OD036435-01	The CFDE Workbench	\$7.2M	15 publications	0 repositories	0 properties
3OT2OD036435-01S2					
3OT2OD036435-01S1					

## Publications

Published works associated with this project.

ID	Title	Authors	RCR	SJR	Citations	Cit./year	Journal	Published	Updated
<a href="#">38643247</a> 	Rummagene: massive mining of gene sets from supporting materials of biomedical research publications	Clarke, Daniel J B ...4 more... Ma'ayan, Avi	5.651	2.071	26	26	Communications biology	2024	Dec 28, 2025
<a href="#">39569206</a> 	RummaGEO: Automatic mining of human and mouse gene sets from GEO.	Marino, Giacomo B ...3 more... Ma'ayan, Avi	2.151	1.527	9	9	Patterns (New York, N.Y.)	2024	Dec 28, 2025
<a href="#">37460679</a> 	Toxicology knowledge graph for structural birth defects.	Evangelista, John Erol ...16 more... Ma'ayan, Avi	2.148	2.098	17	8.5	Communications medicine	2023	Dec 28, 2025
<a href="#">39415993</a> 	The molecular chronology of mammary epithelial cell fate switching.	Vallmajó-Martin, Queralt	0.4	0	2	2	bioRxiv : the preprint server for biology	2024	Dec 28, 2025

<a href="#">38339304</a>	Molecular Atlas of HER2+ Breast Cancer Cells Treated with Endogenous Ligands: Temporal Insights i...	...7 more...	Lytle, Nikki K	Mukund, Kavitha	...10 more...	0.31	145.00	222 Cancers	2024 Dec 28, 2025	
<a href="#">38645198</a>	RummaGEO: Automatic Mining of Human and Mouse Gene Sets from GEO.	Marino, Giacomo B ...2 more...	Subramani am, Shankar	0.24	0	1	1	bioRxiv : the preprint server for biology	2024 Dec 28, 2025	
<a href="#">37953774</a>	Dex-Benchmark: datasets and code to evaluate algorithms for transcriptomics data analysis.	Xie, Zhuorui ...1 more...	Ma'ayan, Avi	0.14	0.625	1	0.5	PeerJ	2023 Dec 28, 2025	
<a href="#">39990419</a>	Spatially distinct cellular and molecular landscapes define prognosis in triple negative breast c...	Mukund, Kavitha	...13 more...	Subramani am, Shankar	0	0	1	1	bioRxiv : the preprint server for biology	2025 Jan 7, 2026

<a href="#">39565691</a>	Protocol for using Multiomics2Targets to identify targets and driver kinases for cancer cohorts p...	Marino, Giacomo B ...6 more... Ma'ayan, Avi	0	0.9 25	0	0	STAR protocols	202 4	Dec 28, 2025
<a href="#">40308216</a>	L2S2: chemical perturbation and CRISPR KO LINCS L1000 signature search engine.	Marino, Giacomo B ...2 more... Ma'ayan, Avi	0	7.7 76	4	4	Nucleic acids research	202 5	Dec 28, 2025

### Publications (cumulative)

Total: 15



### Notes

RCR [Relative Citation Ratio](#)

SJR [Scimago Journal Rank](#)

## </> Repositories

Software repositories associated with this project.

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Description

		S	F	W	C		I	P	R	C	O	C	n	o	L	C	L
T	Last Commit	t	o	a	o		s	R	e	d	t	c	i	o	L	o	a
a		a	r	tc	m		u	A	a	b	r	o	i	t	n	t	g
g		r	k	h	m		e	v	m	u	c	s	e	r	a	g	
s		s	s	e	it		A	g	m	u	c	s	e	b	g	e	
		s	s	rs	s		v	g	e	t	o	b	i	n	.	s	
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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.

Issue/PR Avg Average time issues/pull requests stay open for before being closed.

## Analytics

Website metrics associated with this project.

### Notes

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.