



## Core Project OT2OD030160













### Details











Projects	Name	Award	Publications	Repositories	Analytics
3OT2OD030160-01S3	The LINCS DCIC Engagement Plan with the CFDE	\$3,420,175.00	14 publications	0 repositories	0 properties
3OT2OD030160-01S4					
3OT2OD030160-01S1					
3OT2OD030160-01S2					
1OT2OD030160-01					
3OT2OD030160-01S5					











### 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
<a href="#">37166973</a>  <a href="#">DOI</a> 	Enrichr-KG: bridging enrichment analysis across multiple libraries.	Evangelista , John Erol ...4 more... Ma'ayan, Avi	14. 45 6	0	91	45.5	Nucleic Acids Res	2023	Sep 4, 2025 (just now)
<a href="#">35524556</a>  <a href="#">DOI</a> 	SigCom LINCS: data and metadata search engine for a million gene expression signatures.	Evangelista , John Erol ...11 more... Ma'ayan, Avi	4.8 19	0	57	19	Nucleic Acids Res	2022	Sep 4, 2025 (just now)
<a href="#">33748796</a>  <a href="#">DOI</a> 	Appyters: Turning Jupyter Notebooks into data-driven web apps.	Clarke, Daniel J B ...22 more... Ma'ayan, Avi	4.8 01	0	80	20	Patterns (N Y)	2021	Sep 4, 2025 (just now)
<a href="#">37771373</a>  <a href="#">DOI</a> 	Systems immunology-based drug repurposing framework to target inflammation in atherosclerosis.	Amadori, Letizia ...23 more... Giannarelli, Chiara	2.9 41	0	19	9.5	Nat Cardiova sc Res	2023	Sep 4, 2025 (just now)
<a href="#">35876555</a>  <a href="#">DOI</a> 	Getting Started with LINCS Datasets and Tools.	Xie, Zhuorui ...13	1.4 51	0	14	4.66 7	Curr Protoc	2022	Sep 4, 2025

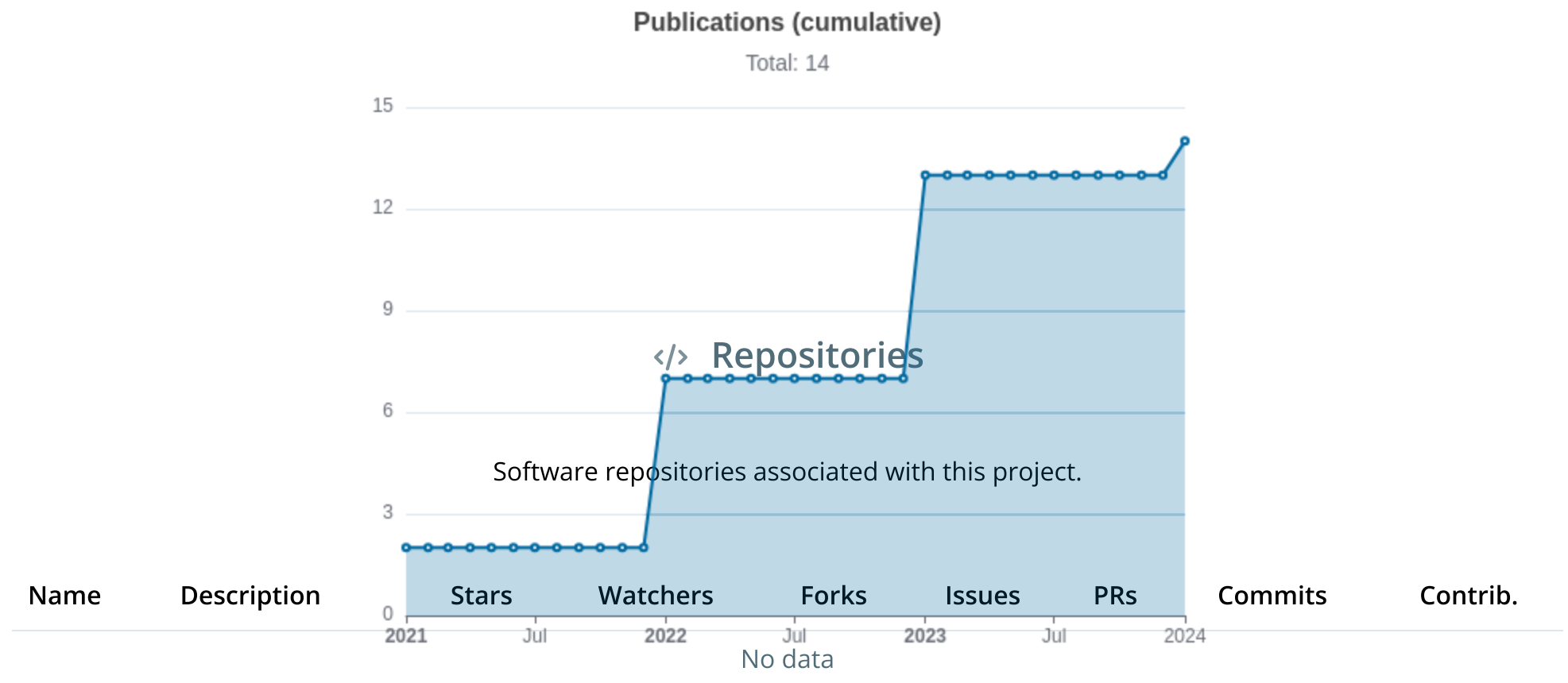
		more... Ma'ayan, Avi								(just now)
<a href="#">35143610</a>  <a href="#">DOI</a> 	blitzGSEA: efficient computation of gene set enrichment analysis through gamma distribution appro...	Lachmann, Alexander ...1 more... Ma'ayan, Avi	1.3 4	0	19	6.33 3	Bioinfor matics	2022	Sep 4, 2025 (just now)	
<a href="#">36869839</a>  <a href="#">DOI</a> 	IncHUB2: aggregated and inferred knowledge about human and mouse lncRNAs.	Marino, Giacomo B ...6 more... Ma'ayan, Avi	1.2 68	0	9	4.5	Databas e (Oxford)	2023	Sep 4, 2025 (just now)	
<a href="#">33787872</a>  <a href="#">DOI</a> 	Drugmonizome and Drugmonizome-ML: integration and abstraction of small molecule attributes for dr...	Kropiwnick i, Eryk ...7 more... Ma'ayan, Avi	1.1 32	0	17	4.25	Databas e (Oxford)	2021	Sep 4, 2025 (just now)	
<a href="#">36874981</a>  <a href="#">DOI</a> 	PrismEXP: gene annotation prediction from stratified gene-gene co-expression matrices.	Lachmann, Alexander ...4 more... Ma'ayan, Avi	0.9 53	0	7	3.5	PeerJ	2023	Sep 4, 2025 (just now)	
<a href="#">36100892</a>  <a href="#">DOI</a> 	Transforming L1000 profiles to RNA-seq-like profiles with deep learning.	Jeon, Minji ...4 more...	0.9 38	0	10	3.33 3	BMC Bioinfor	2022	Sep 4,	

		Ma'ayan, Avi					matrics		2025 (just now)
<a href="#">36409836</a>  <a href="#">DOI</a> 	Making Common Fund data more findable: catalyzing a data ecosystem.	Charbonneau, Amanda L ...40 more... White, Owen	0.818	0	11	3.667	Gigascience	2022	Sep 4, 2025 (just now)
<a href="#">37166966</a>  <a href="#">DOI</a> 	GeneRanger and TargetRanger: processed gene and protein expression levels across cells and tissue...	Marino, Giacomo B ...6 more... Ma'ayan, Avi	0.812	0	6	3	Nucleic Acids Res	2023	Sep 4, 2025 (just now)
<a href="#">37082798</a>  <a href="#">DOI</a> 	Computational screen to identify potential targets for immunotherapeutic identification and remov...	Deng, Eden Z ...4 more... Ma'ayan, Avi	0.732	0	7	3.5	Aging Cell	2023	Sep 4, 2025 (just now)
<a href="#">39127042</a>  <a href="#">DOI</a> 	Multomics2Targets identifies targets from cancer cohorts profiled with transcriptomics, proteomi...	Deng, Eden Z ...6 more... Ma'ayan, Avi	0	0	4	4	Cell Rep Methods	2024	Sep 4, 2025 (just now)

## 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.

# 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 [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.