

# **L** Core Project OT2OD030160

## O Details

Projects	Name	Award	Publications	Repositories	Analytics
	The LINCS DCIC Engagement Plan with the CFDE	\$3,420,175.00	14 publications	0 repositories	0 properties

# 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
<u>37166973</u>	Enrichr-KG: bridging enrichment analysis across multiple libraries.	Evangelista , John Erol 4 more Ma'ayan, Avi	14. 12 1	0	84	42	Nucleic Acids Res	2023	Aug 12, 2025 (just now)
35524556 <b>♂</b> DOI <b>♂</b>	SigCom LINCS: data and metadata search engine for a million gene expression signatures.	Evangelista , John Erol 11 more Ma'ayan, Avi	4.8 79	0	55	18.3 33	Nucleic Acids Res	2022	Aug 12, 2025 (just now)
33748796 <b>♂</b> DOI <b>♂</b>	Appyters: Turning Jupyter Notebooks into data-driven web apps.	Clarke, Daniel J B 22 more Ma'ayan, Avi	4.6 7	0	76	19	Patterns (N Y)	2021	Aug 12, 2025 (just now)
37771373 <b>♂</b> DOI <b>♂</b>	Systems immunology-based drug repurposing framework to target inflammation in atherosclerosis.	Amadori, Letizia 23 more Giannarelli , Chiara	3.0 87	0	19	9.5	Nat Cardiova sc Res	2023	Aug 12, 2025 (just now)
35876555 <b>亿</b> DOI <b>亿</b>	Getting Started with LINCS Datasets and Tools.	Xie, Zhuorui 13	1.4 96	0	14	4.66 7	Curr Protoc	2022	Aug 12, 2025

		more Ma'ayan, Avi							(just now)
36869839 ♂ DOI ♂	IncHUB2: aggregated and inferred knowledge about human and mouse IncRNAs.	Marino, Giacomo B 6 more Ma'ayan, Avi	1.3 31	0	9	4.5	Databas e (Oxford)	2023	Aug 12, 2025 (just now)
35143610 🗗 DOI 🗗	blitzGSEA: efficient computation of gene set enrichment analysis through gamma distribution appro	Lachmann, Alexander 1 more Ma'ayan, Avi	1.2 96	0	18	6	Bioinfor matics	2022	Aug 12, 2025 (just now)
<u>33787872</u> ♂ DOI ♂	Drugmonizome and Drugmonizome-ML: integration and abstraction of small molecule attributes for dr	Kropiwnick i, Eryk 7 more Ma'ayan, Avi	1.0 91	0	16	4	Databas e (Oxford)	2021	Aug 12, 2025 (just now)
36874981 <b>♂</b> DOI <b>♂</b>	PrismEXP: gene annotation prediction from stratified gene-gene co-expression matrices.	Lachmann, Alexander 4 more Ma'ayan, Avi	1	0	7	3.5	PeerJ	2023	Aug 12, 2025 (just now)
36100892 ♂ DOI ♂	Transforming L1000 profiles to RNA-seq-like profiles with deep learning.	Jeon, Minji 4 more	0.9 66	0	10	3.33	BMC Bioinfor	2022	Aug 12,

		Ma'ayan, Avi					matics		2025 (just now)
37166966 🖸	GeneRanger and TargetRanger: processed gene and protein expression levels across cells and tissue	Marino, Giacomo B 6 more Ma'ayan, Avi	0.8 53	0	6	3	Nucleic Acids Res	2023	Aug 12, 2025 (just now)
36409836 <b>♂</b> DOI <b>♂</b>	Making Common Fund data more findable: catalyzing a data ecosystem.	Charbonne au, Amanda L 40 more White, Owen	0.8 41	0	11	3.66 7	Gigascie nce	2022	Aug 12, 2025 (just now)
37082798 🗗 DOI 🗗	Computational screen to identify potential targets for immunotherapeutic identification and remov	Deng, Eden Z 4 more Ma'ayan, Avi	0.7 68	0	7	3.5	Aging Cell	2023	Aug 12, 2025 (just now)
39127042 🗗 DOI 🗗	Multiomics2Targets identifies targets from cancer cohorts profiled with transcriptomics, proteomi	Deng, Eden Z 6 more Ma'ayan, Avi	0	0	2	2	Cell Rep Methods	2024	Aug 12, 2025 (just now)

### Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank





#### 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> 2.

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

Engaged Sessions Visits that had significant interaction <a>\mathbb{Z}</a>.

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