

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> ♂ DOI ♂	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 14, 2025 (just now)
<u>35524556</u> ♂ DOI ♂	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 14, 2025 (just now)
<u>33748796</u> ♂ <u>DOI</u> ♂	Appyters: Turning Jupyter Notebooks into data-driven web apps.	Clarke, Daniel J B22 more Ma'ayan, Avi	4.8 01	0	80	20	Patterns (N Y)	2021	Sep 14, 2025 (just now)
37771373 🗗 DOI 🗗	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 14, 2025 (just now)
35876555 🗹	Getting Started with LINCS Datasets and Tools.	Xie, Zhuorui 13	1.4 51	0	14	4.66 7	Curr Protoc	2022	Sep 14, 2025

		more Ma'ayan, Avi							(just now)
35143610 🗹 DOI 🖸	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 14, 2025 (just now)
36869839 乙	IncHUB2: aggregated and inferred knowledge about human and mouse IncRNAs.	Marino, Giacomo B 6 more Ma'ayan, Avi	1.2 68	0	9	4.5	Databas e (Oxford)	2023	Sep 14, 2025 (just now)
<u>33787872</u> ♂ <u>DOI</u> ♂	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 14, 2025 (just now)
36874981 ♂ DOI ♂	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 14, 2025 (just now)
36100892 亿 <u>DOI</u> 亿	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 14,

		Ma'ayan, Avi					matics		2025 (just now)
36409836 乙	Making Common Fund data more findable: catalyzing a data ecosystem.	Charbonne au, Amanda L 40 more White, Owen	0.8 18	0	11	3.66 7	Gigascie nce	2022	Sep 14, 2025 (just now)
37166966 ☑ DOI ☑	GeneRanger and TargetRanger: processed gene and protein expression levels across cells and tissue	Marino, Giacomo B 6 more Ma'ayan, Avi	0.8 12	0	6	3	Nucleic Acids Res	2023	Sep 14, 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 32	0	7	3.5	Aging Cell	2023	Sep 14, 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	4	4	Cell Rep Methods	2024	Sep 14, 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}.

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