

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 13, 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 13, 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 13, 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 13, 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 13, 2025 |

| | | more Ma'ayan, Avi | | | | | | | (just now) |
|----------------------------|---------------------------------------------------------------------------------------------------------|-----------------------------------------------------|-----------|---|----|-----------|--------------------------|------|-------------------------------------|
| 35143610 🗗 | 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 13, 2025 (just now) |
| 36869839 🗗 DOI 🗗 | 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 13, 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 | 1.1 32 | 0 | 17 | 4.25 | Databas e (Oxford) | 2021 | Sep 13, 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 13, 2025 (just now) |
| 36100892 🗹 DOI 🗹 | 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 13, |

| | | 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 13, 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 13, 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 13, 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 13, 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.