

L Core Project OT2OD030160

O Details

| Projects | Name | Award | Publications | Repositories | Analytics |
|--|--|----------------|-----------------|----------------|--------------|
| 1OT2OD030160-01 3OT2OD030160-01S2 3OT2OD030160-01S5 3OT2OD030160-01S3 3OT2OD030160-01S1 3OT2OD030160-01S4 | 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 | R C R | SJ R | Cita tion s | Cit./ yea r | Journal | Publ ishe d | Upda ted |
|---------------------|--|--|-------------|---------------|-------------------|-------------------|---------------------------------------|-------------------|---|
| 37166973 🗗 DOI 🗗 | Enrichr-KG: bridging enrichment analysis across multiple libraries. | John Erol Evangelist a 4 more Avi Ma'ayan | 7. 82 | 7. 04 8 | 24 | 24 | Nucleic Acids Research | 2,02 3 | Sep 28, 2024 (2 weeks ago) |
| 33748796 🖸 DOI 🗗 | Appyters: Turning Jupyter Notebooks into data-driven web apps. | Daniel J B Clarke 22 more Avi Ma'ayan | 4. 58 | 0 | 55 | 18. 333 | Patterns (N Y) | 2,02 1 | Sep 28, 2024 (2 weeks ago) |
| 35524556 🗹 DOI 🗹 | SigCom LINCS: data and metadata search engine for a million gene expression signatures. | John Erol Evangelist a 11 more Avi Ma'ayan | 4. 19 | 7. 04 8 | 30 | 15 | Nucleic Acids Research | 2,02 2 | Sep 28, 2024 (2 weeks ago) |
| 37771373 🗗 DOI 🗗 | Systems immunology-based drug repurposing framework to target inflammation in atherosclerosis. | Letizia Amadori 23 more Chiara Giannarell | 2. 73 | 3. 71 | 8 | 8 | Nature Cardiovascul ar Research | 2,02 3 | Sep 28, 2024 (2 weeks ago) |

| 33787872 🗗 DOI 🗗 | Drugmonizome and Drugmonizome-ML: integration and abstraction of small molecule attributes for dr | Eryk Kropiwnic ki 7 more Avi Ma'ayan | 1. 35 | 0 | 14 | 4.6 67 | Database (Oxford) | 2,02 1 | Sep 28, 2024 (2 weeks ago) |
|---------------------|---|--|----------|---------------|----|-----------|---------------------------|-----------|---|
| 35143610 🗹 DOI 🗹 | blitzGSEA: efficient computation of gene set enrichment analysis through gamma distribution appro | Alexander Lachmann 1 more Avi Ma'ayan | 1. 02 | 2. 57 4 | 9 | 4.5 | Bioinformatic s | 2,02 2 | Sep 28, 2024 (2 weeks ago) |
| 36409836 ☑ DOI ☑ | Making Common Fund data more findable: catalyzing a data ecosystem. | Amanda L Charbonn eau 40 more Owen White | 0. 91 | 4. 62 1 | 8 | 4 | GigaScience | 2,02 2 | Sep 28, 2024 (2 weeks ago) |
| 36100892 🗗 DOI 🗗 | Transforming L1000 profiles to RNA-seq- like profiles with deep learning. | Minji Jeon 4 more Avi Ma'ayan | 0. 61 | 1. 00 5 | 4 | 2 | BMC Bioinformatic s | 2,02 2 | Sep 28, 2024 (2 weeks ago) |

| 35876555 🗹 DOI 🗹 | Getting Started with LINCS Datasets and Tools. | Zhuorui Xie 13 more Avi Ma'ayan | 0. 59 | 1. 52 | 3 | 1.5 | Current Protocols | 2,02 2 | Sep 28, 2024 (2 weeks ago) |
|-----------------------------------|---|---|----------|---------------|---|-----|---------------------------|-----------|---|
| 37166966 ☑ DOI ☑ | GeneRanger and TargetRanger: processed gene and protein expression levels across cells and tissue | Giacomo B Marino 6 more Avi Ma'ayan | 0 | 7. 04 8 | 2 | 2 | Nucleic Acids Research | 2,02 3 | Sep 28, 2024 (2 weeks ago) |
| 36869839 ♂ DOI ♂ | IncHUB2: aggregated and inferred knowledge about human and mouse IncRNAs. | Giacomo B Marino 6 more Avi Ma'ayan | 0 | 0 | 3 | 3 | Database (Oxford) | 2,02 3 | Sep 28, 2024 (2 weeks ago) |
| 37082798 🗷 DOI 🗹 | Computational screen to identify potential targets for immunotherapeutic identification and remov | Eden Z Deng 4 more Avi Ma'ayan | 0 | 2. 90 4 | 3 | 3 | Aging Cell | 2,02 3 | Sep 28, 2024 (2 weeks ago) |

| 36874981 🖸 DOI 🗗 | PrismEXP: gene annotation prediction from stratified gene-gene co-expression matrices. | Alexander Lachmann 4 more Avi Ma'ayan | 0 | 0. 62 3 | 4 | 4 | PeerJ | 2,02 3 | Sep 28, 2024 (2 weeks ago) |
|---------------------|---|---|---|---------------|---|---|-------------------------|-----------|---|
| 39127042 🗗 DOI 🗗 | Multiomics2Targets identifies targets from cancer cohorts profiled with transcriptomics, proteomi | Eden Z Deng 6 more Avi Ma'ayan | 0 | 2. 10 7 | 0 | 0 | Cell Reports Methods | 2,02 4 | Sep 28, 2024 (2 weeks ago) |

Notes

RCR Relative Citation Ratio

SJR Scimago Journal Rank

</> Repositories

Software repositories associated with this project.

| Name | Description | Stars | Wateher | Slative Publications Is | sues Pf | Rs | Commits | Contrib. |
|--------------|-----------------------|-------------------------|-------------|-----------------------------|-----------|-------|------------|--------------|
| | 45 | | | Total: 14 No data | | | | |
| | 15 — | | | | | | | |
| Name Tag | gs Last Commit — | Avg Issue | Avg PR | Languages Licens | se Readme | e Con | ntributing | Dependencies |
| | | | | No data | | | | |
| | 9 — | | | | | | | |
| Notes | 6 — | | | | | | | |
| Repository | For storing, tracking | g changes to | , and colla | borating on a piece of | software. | | | |
| PR | "Pull request", a dra | it change (n | ew feature | e, bug fix, etc.) to a rep | 0. | | | |
| Closed/Open | Resolved/unresolve | ed. | | | | | | |
| Avg Issue/PR | Average time is sugg | s/pull reques | sts stay/op | en for before bein့g္တင္ခါ၀ | osed. | 2024 | | |

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> **2**.

Engaged Sessions <u>Visits that had significant interaction</u> **?**.

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

Generated on Oct 10, 2024

Developed with support from NIH Award U54 OD036472