



Core Project OT2OD036435



Details











| Projects | Name | Award | Publications | Repositories | Analytics |
|--------------------------------------|--------------------|----------------|-----------------|----------------|--------------|
| 1OT2OD036435-01 3OT2OD036435-01S1 | The CFDE Workbench | \$3,250,000.00 | 13 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 |
|---|--|---------------------------------------|-----------|---------|-------------------|-------------------|----------------|-------------------|--------------------|
| 38643247 DOI | Rummagene: massive mining of gene sets from supporting materials of biomedical research publicati... | Clarke, Daniel J B ...4 more... | 5.0 47 | 0 | 16 | 16 | Commun Biol | 2024 | Aug 30, 2025 |

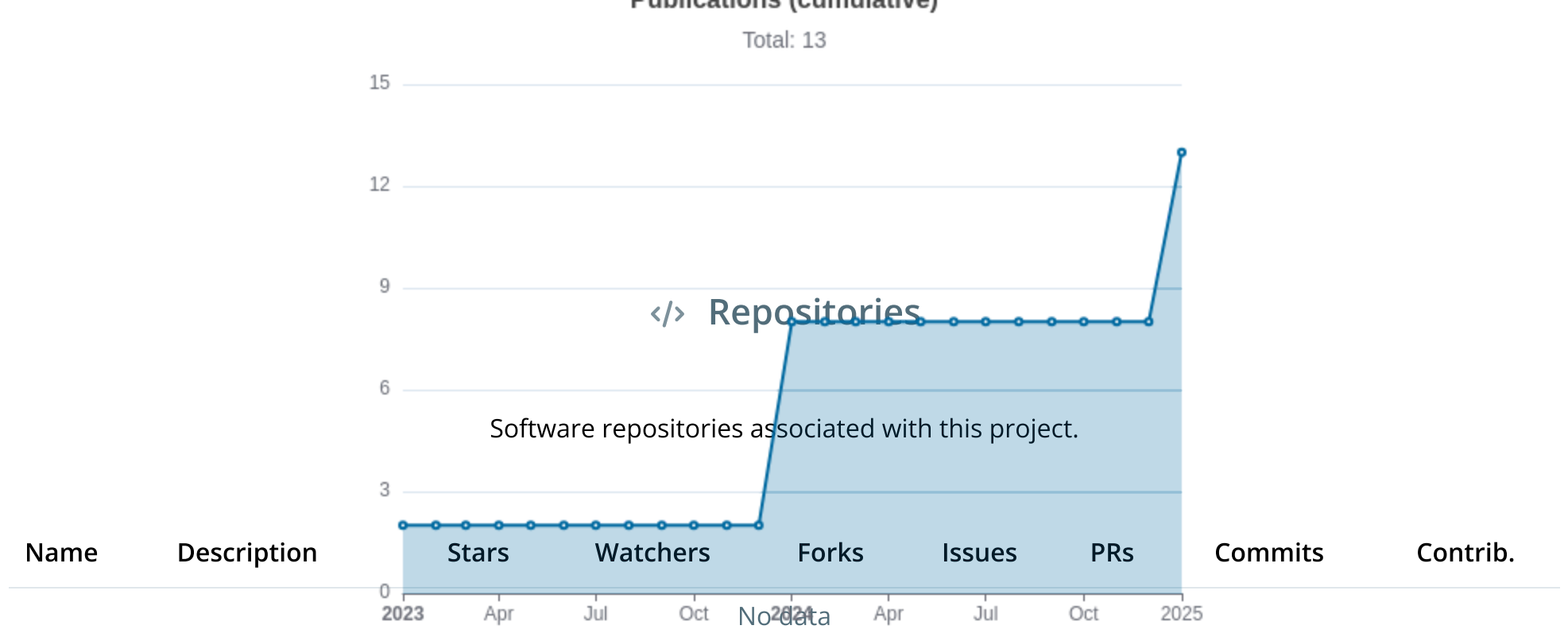
| | | | | | | | | | | |
|---|--|---|-------|---|----|-----|-------------------|------|-------------------------|------------|
| | | Ma'ayan, Avi | | | | | | | | (just now) |
| 39569206  DOI  | RummaGEO: Automatic mining of human and mouse gene sets from GEO. | Marino, Giacomo B ...3 more... Ma'ayan, Avi | 2.76 | 0 | 8 | 8 | Patterns (N Y) | 2024 | Aug 30, 2025 (just now) | |
| 37460679  DOI  | Toxicology knowledge graph for structural birth defects. | Evangelista , John Erol ...16 more... Ma'ayan, Avi | 2.122 | 0 | 13 | 6.5 | Commun Med (Lond) | 2023 | Aug 30, 2025 (just now) | |
| 37953774  DOI  | Dex-Benchmark: datasets and code to evaluate algorithms for transcriptomics data analysis. | Xie, Zhuorui ...1 more... Ma'ayan, Avi | 0.188 | 0 | 1 | 0.5 | PeerJ | 2023 | Aug 30, 2025 (just now) | |
| 39565691  DOI  | Protocol for using Multiomics2Targets to identify targets and driver kinases for cancer cohorts p... | Marino, Giacomo B ...6 more... Ma'ayan, Avi | 0 | 0 | 0 | 0 | STAR Protoc | 2024 | Aug 30, 2025 (just now) | |
| 40308216  DOI  | L2S2: chemical perturbation and CRISPR KO LINCS L1000 signature search engine. | Marino, Giacomo B | 0 | 0 | 1 | 1 | Nucleic Acids Res | 2025 | Aug 30, | |

| | | | | | | | | | | |
|---|--|---|---|---|---|---|------------|------|-------------------------------------|-----------------------|
| | | ...2 more... Ma'ayan, Avi | | | | | | | | 2025 (just now) |
| 40133794  DOI  | IncRNAlyzr: Enrichment Analysis for IncRNA Sets. | Evangelista , John Erol ...5 more... Ma'ayan, Avi | 0 | 0 | 2 | 2 | J Mol Biol | 2025 | Aug 30, 2025 (just now) | |
| 40672200  DOI  | A novel subtyping method for TNBC with implications for prognosis and therapy. | Mesrizade h, Zahra ...9 more... Subramani am, Shankar | 0 | 0 | 0 | 0 | bioRxiv | 2025 | Aug 30, 2025 (just now) | |
| 39990419  DOI  | Spatially distinct cellular and molecular landscapes define prognosis in triple negative breast c... | Mukund, Kavitha ...13 more... Subramani am, Shankar | 0 | 0 | 0 | 0 | bioRxiv | 2025 | Aug 30, 2025 (just now) | |
| 38645198  DOI  | RummaGEO: Automatic Mining of Human and Mouse Gene Sets from GEO. | Marino, Giacomo B ...2 more... Ma'ayan, Avi | 0 | 0 | 1 | 1 | bioRxiv | 2024 | Aug 30, 2025 (just now) | |

| | | | | | | | | | |
|--|--|--|---|---|---|---|-----------------|------|-------------------------|
| 38339304  DOI  | Molecular Atlas of HER2+ Breast Cancer Cells Treated with Endogenous Ligands: Temporal Insights i... | Mukund, Kavitha ...10 more... Subramani am, Shankar | 0 | 0 | 1 | 1 | Cancers (Basel) | 2024 | Aug 30, 2025 (just now) |
| 39415993  DOI  | The molecular chronology of mammary epithelial cell fate switching. | Vallmajo-Martin, Queralt ...7 more... Lytle, Nikki K | 0 | 0 | 0 | 0 | bioRxiv | 2024 | Aug 30, 2025 (just now) |
| 40501705  DOI  | A Gene Set Foundation Model Pre-Trained on a Massive Collection of Diverse Gene Sets. | Clarke, Daniel J B ...1 more... Ma'ayan, Avi | 0 | 0 | 0 | 0 | bioRxiv | 2025 | Aug 30, 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.

Built on Aug 30, 2025

Developed with support from NIH Award [U54 OD036472](#)