



## (Core Project OT2OD036435)

### ◎ Details

| Projects          | Name               | Award          | Publications    | Repositories   | Analytics    |
|-------------------|--------------------|----------------|-----------------|----------------|--------------|
| 1OT2OD036435-01   | The CFDE Workbench | \$7,184,710.00 | 15 publications | 0 repositories | 0 properties |
| 3OT2OD036435-01S2 |                    |                |                 |                |              |
| 3OT2OD036435-01S1 |                    |                |                 |                |              |

### ❑ Publications

Published works associated with this project.

| ID                       | Title  | Authors            | RC<br>R   | SJ<br>R | Cita<br>tion<br>s | Cit./<br>yea<br>r | Journal     | Publ<br>ishe<br>d | Upda<br>ted |
|--------------------------|--|--------------------|-----------|---------|-------------------|-------------------|-------------|-------------------|-------------|
| <a href="#">38643247</a> | Rummagene: massive mining of gene sets from supporting materials of biomedical | Clarke, Daniel J B | 4.9<br>09 | 0       | 18                | 18                | Commun Biol | 2024              | Oct 15,     |



|                          |  |   |         |             |      |                         |
|--------------------------|--|---|---------|-------------|------|-------------------------|
| <a href="#">39415993</a> | The molecular chronology of mammary epithelial cell fate switching.                                  | Vallmajó-Martin,<br>Queralt<br>...7 more...<br>Lytle, Nikki K | 0 0 0 0 | bioRxiv     | 2024 | Oct 15, 2025 (just now) |
| <a href="#">40133794</a> | lncRNAlzr: Enrichment Analysis for lncRNA Sets.  | Evangelista, John Erol<br>...5 more...<br>Ma'ayan, Avi        | 0 0 3 3 | J Mol Biol  | 2025 | Oct 15, 2025 (just now) |
| <a href="#">40832173</a> | ChEA-KG: Human Transcription Factor Regulatory Network with a Knowledge Graph Interactive User In... | Byrd, Anna I<br>...4 more...<br>Ma'ayan, Avi                  | 0 0 0 0 | bioRxiv     | 2025 | Oct 15, 2025 (just now) |
| <a href="#">39565691</a> | Protocol for using Multiomics2Targets to identify targets and driver kinases for cancer cohorts p... | Marino, Giacomo B<br>...6 more...<br>Ma'ayan, Avi             | 0 0 0 0 | STAR Protoc | 2024 | Oct 15, 2025 (just now) |

|                          |  |                                     |         |                   |      |                         |
|--------------------------|--|-------------------------------------|---------|-------------------|------|-------------------------|
| <a href="#">40956607</a> | Creating an Interactive Web Interface for Networks Stored in Knowledge Graph Databases.              | Evangelista, John Erol ...5 more... | 0 0 1 1 | Curr Protoc       | 2025 | Oct 15, 2025 (just now) |
| <a href="#">40672200</a> | A novel subtyping method for TNBC with implications for prognosis and therapy.                       | Mesrizadeh, Zahra ...9 more...      | 0 0 0 0 | bioRxiv           | 2025 | Oct 15, 2025 (just now) |
| <a href="#">39990419</a> | Spatially distinct cellular and molecular landscapes define prognosis in triple negative breast c... | Mukund, Kavitha ...13 more...       | 0 0 1 1 | bioRxiv           | 2025 | Oct 15, 2025 (just now) |
| <a href="#">40308216</a> | L2S2: chemical perturbation and CRISPR KO LINCS L1000 signature search engine.                       | Marino, Giacomo B ...2 more...      | 0 0 1 1 | Nucleic Acids Res | 2025 | Oct 15, 2025 (just now) |

|                          |  |   |   |   |   |   |                    |      |                         |
|--------------------------|--|---|---|---|---|---|--------------------|------|-------------------------|
| <a href="#">38339304</a> | Molecular Atlas of HER2+ Breast Cancer Cells Treated with Endogenous Ligands: Temporal Insights i... | Mukund,<br>Kavitha<br>...10<br>more...<br>Subramani<br>am,<br>Shankar | 0 | 0 | 1 | 1 | Cancers<br>(Basel) | 2024 | Oct 15, 2025 (just now) |
| <a href="#">40501705</a> | A Gene Set Foundation Model Pre-Trained on a Massive Collection of Diverse Gene Sets.                | Clarke,<br>Daniel J B<br>...1 more...<br>Ma'ayan,<br>Avi              | 0 | 0 | 0 | 0 | bioRxiv            | 2025 | Oct 15, 2025 (just now) |

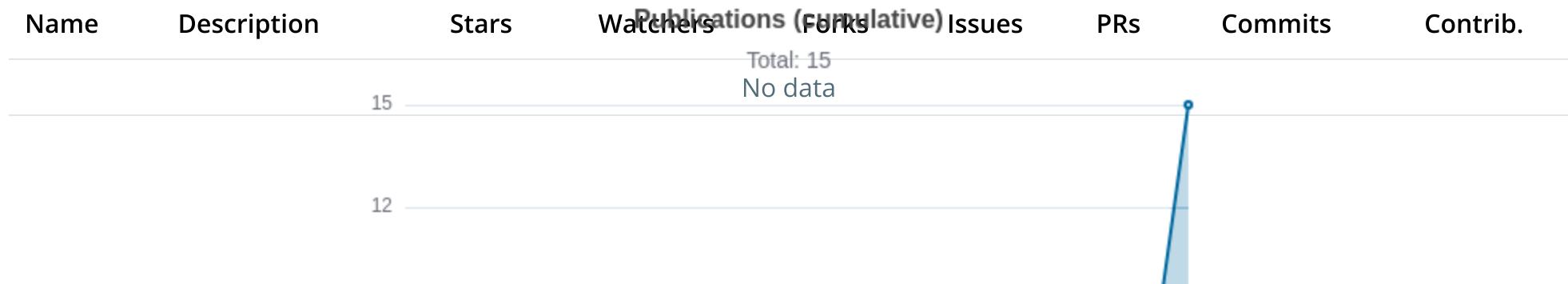
## Notes

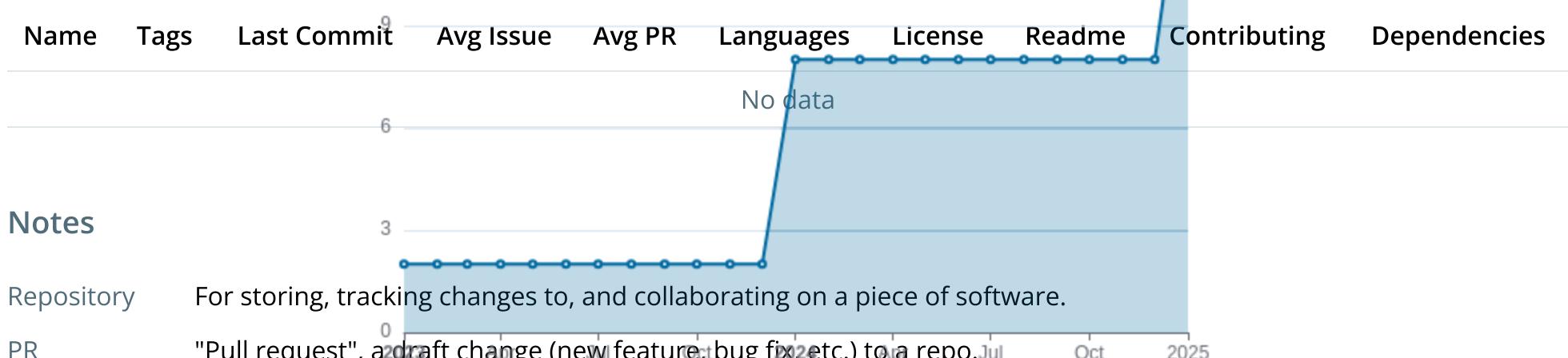
### getRepository

RCR [Relative Citation Ratio](#)

SJR [Scimago Journal Rank](#)

Software repositories associated with this project.





**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. Jul 2018

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.

# 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.

