



(Core Project R03OD039975)

Overview

High-level info about this project.

| Projects | Name | Award | Publications | Repositories | Analytics |
|-----------------|--|--------|----------------|----------------|--------------|
| 1R03OD039975-01 | Rare variant analysis integrating large public DNA sequencing controls and gene expression data for discovering novel predisposition genes of orofacial clefts | \$321K | 0 publications | 0 repositories | 0 properties |

Publications

Published works associated with this project.

| ID | Title | Authors | RCR | SJR | Citations | Cit./year | Journal | Published | Updated |
|---------|-------|---------|-----|-----|-----------|-----------|---------|-----------|---------|
| No data | | | | | | | | | |

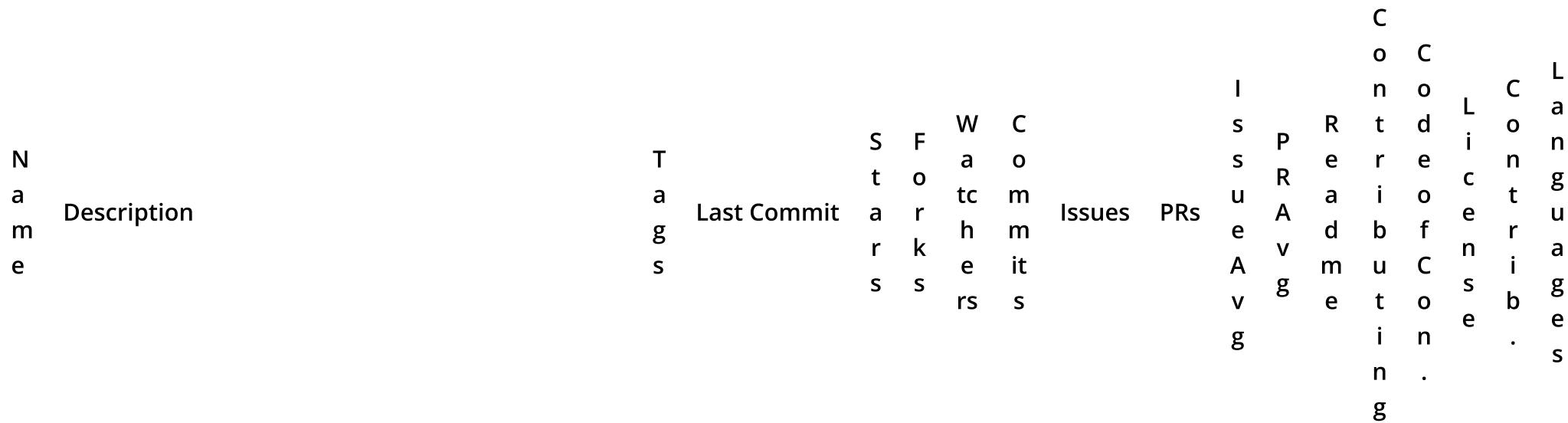
Notes

RCR [Relative Citation Ratio](#) 

SJR [Scimago Journal Rank](#) 

</> Repositories

Software repositories associated with this project.



No data

Built on Jan 29, 2026

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

PR "Pull request", a draft change (new feature, bug fix, etc.) to a repo.

Closed/Open Resolved/unresolved.

Issue/PR Avg 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

Website metrics 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.