

# Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

## CaltechDATA

RRID:SCR\_017602

Type: Tool

### Proper Citation

CaltechDATA (RRID:SCR\_017602)

### Resource Information

**URL:** <https://data.caltech.edu/>

**Proper Citation:** CaltechDATA (RRID:SCR\_017602)

**Description:** Data and software repository from CalTech.

**Resource Type:** data repository, service resource, software repository, software resource, storage service resource

**Keywords:** Data, dataset, respository, CalTech

**Availability:** Restricted

**Resource Name:** CaltechDATA

**Resource ID:** SCR\_017602

**Alternate IDs:** DOI:10.17616/R3sw99, r3d100012384, DOI:10.22002, DOI:10.25504/FAIRsharing.S09se7

**Alternate URLs:** <https://doi.org/10.17616/r3sw99>, <https://doi.org/10.22002/>, <https://dx.doi.org/10.22002/>, <https://fairsharing.org/10.25504/FAIRsharing.S09se7>, <https://doi.org/10.17616/R3SW99>

**Record Creation Time:** 20220129T080336+0000

**Record Last Update:** 20260905T132826+0000

### Ratings and Alerts

No rating or validation information has been found for CaltechDATA.

No alerts have been found for CaltechDATA.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 11 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

Chadly D, et al. (2026) Regenerative base editing enables deep lineage recording. bioRxiv : the preprint server for biology.

Ouadah Y, et al. (2026) Evolutionary basis of male same-sex sexual behavior by multiple pheromone switches in Drosophila. Current biology : CB, 36(7), 1716.

Fan C, et al. (2026) Bivalent mRNA booster encoding virus-like particles elicits potent polyclonal receptor-binding domain antibodies in pre-vaccinated mice. eLife, 14.

Kuintzle R, et al. (2025) Diversity in Notch ligand-receptor signaling interactions. eLife, 12.

Du R, et al. (2024) miRNA circuit modules for precise, tunable control of gene expression. bioRxiv : the preprint server for biology.

Xia S, et al. (2024) Synthetic protein circuits for programmable control of mammalian cell death. Cell, 187(11), 2785.

Briney KA, et al. (2024) Measuring data rot: An analysis of the continued availability of shared data from a Single University. PloS one, 19(6), e0304781.

Kuintzle R, et al. (2023) Diversity in Notch ligand-receptor signaling interactions. bioRxiv : the preprint server for biology.

da Veiga Beltrame E, et al. (2022) WormBase single-cell tools. Bioinformatics advances, 2(1), vbac018.

Segalin C, et al. (2021) The Mouse Action Recognition System (MARS) software pipeline for automated analysis of social behaviors in mice. eLife, 10.

B??guin JB, et al. (2020) Reduced volume and reflection for bright optical tweezers with radial Laguerre-Gauss beams. Proceedings of the National Academy of Sciences of the United States of America, 117(42), 26109.