

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 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 [Kravitz](#) .

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.

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

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.

Bergouin 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.