

# Resource Summary Report

Generated by [kravitz2](#) on Jul 30, 2026

## [Trim28<sup>MommeD9</sup>/Trim28<sup>+</sup>](#)

RRID:MGI:3821610

Type: Organism

### Proper Citation

RRID:MGI:3821610

### Organism Information

**URL:**

**Proper Citation:** RRID:MGI:3821610

**Description:** Allele Detail: Chemically induced (ENU) This is a legacy resource.

**Species:** Mus musculus

**Notes:** Allele Detail: Chemically induced (ENU) This is a legacy resource.

**Phenotype:** decreased litter size, abnormal sex determination

**Affected Gene:** Trim28

**Genomic Alteration:** MommeD9

**Catalog Number:** 3821610

**Background:** involves: FVB/N

**Database:** MGI, Mouse Genome Informatics MGI

**Database Abbreviation:** MGI

**Availability:** Availability unknown check source stock center

**Source References:** [PMID:19099580](#)

**Organism Name:** Trim28<sup>MommeD9</sup>/Trim28<sup>+</sup>

**Record Creation Time:** 20240120T190403+0000

**Record Last Update:** 20240130T201858+0000

### Ratings and Alerts

No rating or validation information has been found for Trim28<sup>MommeD9</sup>/Trim28<sup>+</sup>.

No alerts have been found for Trim28<sup>MommeD9</sup>/Trim28<sup>+</sup>.

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## Data and Source Information

**Source:** [Integrated Animals](#)

**Source Database:** MGI, Mouse Genome Informatics MGI

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Costello KR, et al. (2021) Sequence features of retrotransposons allow for epigenetic variability. eLife, 10.