

Resource Summary Report

Generated by [NIF](#) on Aug 4, 2026

[Trim28^{MommeD9}/Trim28⁺](#)

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^{MommeD9}/Trim28⁺

Record Creation Time: 20240120T190403+0000

Record Last Update: 20240130T201858+0000

Ratings and Alerts

No rating or validation information has been found for Trim28^{MommeD9}/Trim28⁺.

No alerts have been found for Trim28^{MommeD9}/Trim28⁺.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: MGI, Mouse Genome Informatics MGI

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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