

Resource Summary Report

Generated by SPARC SAWG on Aug 8, 2026

Adult Mouse Anatomy Ontology

RRID:SCR_006568

Type: Tool

Proper Citation

Adult Mouse Anatomy Ontology (RRID:SCR_006568)

Resource Information

URL: http://www.informatics.jax.org/searches/AMA_form.shtml

Proper Citation: Adult Mouse Anatomy Ontology (RRID:SCR_006568)

Description: Ontology that organizes anatomical structures for the adult mouse (Theiler stage 28) spatially and functionally, using "is a" and "part of" relationships. The ontology is used to describe expression data for the adult mouse and phenotype data pertinent to anatomy in standardized ways. The browser can be used to view anatomical terms and their relationships in a hierarchical display.

Abbreviations: MA

Synonyms: Adult Mouse Anatomy Browser, MGI Adult Mouse Anatomical Dictionary Browser, Adult Mouse Anatomical Dictionary Browser, Mouse Adult Gross Anatomy Ontology, Anatomical Dictionary for the Adult Mouse

Resource Type: ontology, data or information resource, controlled vocabulary

Keywords: functionally, adult mouse, anatomical, anatomy, phenotype, postnatal, structure, theiler stage 28, obo, gene expression

Funding: NIH ;
NICHD HD33745;
NICHD F32 HD08435-01;
NHGRI F32 HG00215-01

Availability: Acknowledgement requested

Resource Name: Adult Mouse Anatomy Ontology

Resource ID: SCR_006568

Alternate IDs: nif-0000-10300

Alternate URLs: <http://purl.bioontology.org/ontology/MA>

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260808T185846+0000

Ratings and Alerts

No rating or validation information has been found for Adult Mouse Anatomy Ontology.

No alerts have been found for Adult Mouse Anatomy Ontology.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Jiao X, et al. (2025) Progressive gut microbiota shifts and functional alterations across aging stages and frailty in mice. iScience, 28(7), 112985.

Xue X, et al. (2021) Matching Biomedical Ontologies through Adaptive Multi-Modal Multi-Objective Evolutionary Algorithm. Biology, 10(12).

West DB, et al. (2015) A lacZ reporter gene expression atlas for 313 adult KOMP mutant mouse lines. Genome research, 25(4), 598.