

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

Functional Annotation

RRID:SCR_017519

Type: Tool

Proper Citation

Functional Annotation (RRID:SCR_017519)

Resource Information

URL: <http://www.informatics.jax.org/function.shtml>

Proper Citation: Functional Annotation (RRID:SCR_017519)

Description: MGI GO project provides functional annotations for mouse gene products using Gene Ontology. Functional annotation using Gene Ontology (GO).

Resource Type: service resource

Keywords: MGI, GO, functional, annotation, mouse, gene, product, Gene Ontology

Availability: Free, Freely available

Resource Name: Functional Annotation

Resource ID: SCR_017519

Record Creation Time: 20220129T080335+0000

Record Last Update: 20260919T195341+0000

Ratings and Alerts

No rating or validation information has been found for Functional Annotation.

No alerts have been found for Functional Annotation.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Dimond A, et al. (2025) PBK/TOPK mediates Ikaros, Aiolos and CTCF displacement from mitotic chromosomes and alters chromatin accessibility at selected C2H2-zinc finger protein binding sites. Nature communications, 16(1), 8348.

Bujanover N, et al. (2021) Hypersensitivity response has negligible impact on Hematopoietic Stem Cells. Stem cell reports, 16(8), 1884.

Sanavia T, et al. (2021) Temporal Transcriptome Analysis Reveals Dynamic Gene Expression Patterns Driving ??-Cell Maturation. Frontiers in cell and developmental biology, 9, 648791.

Lu T, et al. (2020) Investigating transcriptome-wide sex dimorphism by multi-level analysis of single-cell RNA sequencing data in ten mouse cell types. Biology of sex differences, 11(1), 61.