Resource Summary Report

Generated by NIF on Apr 26, 2025

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

Funding:

Availability: Free, Freely available

Resource Name: Functional Annotation

Resource ID: SCR_017519

Record Creation Time: 20220129T080335+0000

Record Last Update: 20250420T014839+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 3 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

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.