

Resource Summary Report

Generated by [NIF](#) on Sep 15, 2026

Mouse Transcriptome Project

RRID:SCR_007311

Type: Tool

Proper Citation

Mouse Transcriptome Project (RRID:SCR_007311)

Resource Information

URL: <http://www.ncbi.nlm.nih.gov/genome/guide/mouse/MouseTranscriptome.html>

Proper Citation: Mouse Transcriptome Project (RRID:SCR_007311)

Description: Focus is the generation of a reference expression database for the C57BL/6J mouse. The current dataset will consist of 90 samples, primarily from adult tissue, and was analyzed using a technique known as Massively Parallel Signature Sequencing (MPSS). The data are stored in the Gene Expression Omnibus (GEO). The focus of this project is the generation of a reference RNA transcriptome expression database for the C57BL/6J mouse. The current dataset will consist of 90 samples, primarily from adult tissue, and was analyzed using a technique known as Massively Parallel Signature Sequencing (MPSS)

Abbreviations: Mouse Transcriptome Project

Synonyms: NIH Mouse Transcriptome Project

Resource Type: data or information resource, database

Funding: NIH Blueprint for Neuroscience Research

Resource Name: Mouse Transcriptome Project

Resource ID: SCR_007311

Alternate IDs: nif-0000-00176

Record Creation Time: 20220129T080241+0000

Record Last Update: 20260912T200146+0000

Ratings and Alerts

No rating or validation information has been found for Mouse Transcriptome Project.

No alerts have been found for Mouse Transcriptome Project.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Galante PA, et al. (2007) Sense-antisense pairs in mammals: functional and evolutionary considerations. Genome biology, 8(3), R40.