

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## MAGE-TAB

RRID:SCR\_003222

Type: Tool

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### Proper Citation

MAGE-TAB (RRID:SCR\_003222)

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### Resource Information

**URL:** <http://www.mged.org/mage-tab/>

**Proper Citation:** MAGE-TAB (RRID:SCR\_003222)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 27,2023. A simple tab-delimited, spreadsheet-based format which will become a part of the MAGE microarray data standard that can be used for annotating and communicating microarray data in a MIAME compliant fashion. MAGE-TAB will enable laboratories without bioinformatics experience or support to manage, exchange and submit well-annotated microarray data in a standard format using a spreadsheet. The MAGE-TAB format is self-contained, and does not require an understanding of MAGE-ML or XML.

**Abbreviations:** MAGE-TAB

**Resource Type:** data or information resource, narrative resource, standard specification

**Defining Citation:** [PMID:17087822](#)

**Keywords:** microarray, bio.tools

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** MAGE-TAB

**Resource ID:** SCR\_003222

**Alternate IDs:** nlx\_157259, biotools:tab2mage

**Alternate URLs:** <https://bio.tools/tab2mage>

**Record Creation Time:** 20220129T080217+0000

**Record Last Update:** 20260905T132459+0000

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### Ratings and Alerts

No rating or validation information has been found for MAGE-TAB.

No alerts have been found for MAGE-TAB.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Skrzypiec AE, et al. (2013) Stress-induced lipocalin-2 controls dendritic spine formation and neuronal activity in the amygdala. PloS one, 8(4), e61046.

Kristoffersen SM, et al. (2012) Global mRNA decay analysis at single nucleotide resolution reveals segmental and positional degradation patterns in a Gram-positive bacterium. Genome biology, 13(4), R30.

Byrne M, et al. (2012) VarioML framework for comprehensive variation data representation and exchange. BMC bioinformatics, 13, 254.

Wang X, et al. (2012) Inactivation of a novel FGF23 regulator, FAM20C, leads to hypophosphatemic rickets in mice. PLoS genetics, 8(5), e1002708.

Washington NL, et al. (2011) The modENCODE Data Coordination Center: lessons in harvesting comprehensive experimental details. Database : the journal of biological databases and curation, 2011, bar023.