

# Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

## GSE13732

RRID:SCR\_003648

Type: Tool

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### Proper Citation

GSE13732 (RRID:SCR\_003648)

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### Resource Information

**URL:** <http://ranchobiosciences.com/gse13732/>

**Proper Citation:** GSE13732 (RRID:SCR\_003648)

**Description:** Curated data set from a study that developed biomarkers that may predict development of Clinically Isolated Syndrome (CIS) into a full multiple sclerosis. Expression data was taken from 37 CIS patients and 28 healthy controls at baseline. 34 CIS patients and 10 healthy controls were resampled at a second time point, approximately one year later. Patients were followed clinically for up to two years to determine the TTC (time to conversion to MS).

**Abbreviations:** GSE13732

**Resource Type:** data set, data or information resource

**Keywords:** expression, adult human

**Related Condition:** Inflammatory disease, Clinically Isolated Syndrome, Multiple Sclerosis, Healthy control

**Availability:** Free, Public

**Resource Name:** GSE13732

**Resource ID:** SCR\_003648

**Alternate IDs:** nlx\_157800

**Record Creation Time:** 20220129T080220+0000

**Record Last Update:** 20260805T044057+0000

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### Ratings and Alerts

No rating or validation information has been found for GSE13732.

No alerts have been found for GSE13732.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Manian M, et al. (2021) An Integrated Bioinformatics Analysis of the Potential Regulatory Effects of miR-21 on T-cell Related Target Genes in Multiple Sclerosis. Avicenna journal of medical biotechnology, 13(3), 149.

Shin SH, et al. (2020) Synthetic lethality by targeting the RUVBL1/2-TTT complex in mTORC1-hyperactive cancer cells. Science advances, 6(31), eaay9131.