

Resource Summary Report

Generated by [NIF](#) on Aug 9, 2026

GSE13732

RRID:SCR_003648

Type: Tool

Proper Citation

GSE13732 (RRID:SCR_003648)

Resource Information

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

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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 or information resource, data set

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: 20260808T190549+0000

Ratings and Alerts

No rating or validation information has been found for GSE13732.

No alerts have been found for GSE13732.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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.