

# Resource Summary Report

Generated by [RRID](#) on Sep 19, 2026

## OMiCC

RRID:SCR\_016604

Type: Tool

### Proper Citation

OMiCC (RRID:SCR\_016604)

### Resource Information

**URL:** <https://omicc.niaid.nih.gov>

**Proper Citation:** OMiCC (RRID:SCR\_016604)

**Description:** Community based, biologist friendly web platform for creating and meta analyzing annotated gene expression data compendia., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

**Abbreviations:** OMiCC

**Synonyms:** OMics Compendia Commons

**Resource Type:** analysis service resource, data analysis service, data or information resource, production service resource, service resource, software resource, web application

**Defining Citation:** [PMID:27323300](#)

**Keywords:** creating, metadata, analysis, annotated, gene, expression, data, compendia, human, mouse

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

**Resource Name:** OMiCC

**Resource ID:** SCR\_016604

**Record Creation Time:** 20220129T080331+0000

**Record Last Update:** 20260919T195516+0000

### Ratings and Alerts

No rating or validation information has been found for OMiCC.

No alerts have been found for OMiCC.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Lima FT, et al. (2025) The Dual Immunoregulatory Role of CREB3L1 Underlying Latent and Severe Tuberculosis Clinical Manifestation. Immunology.

Liu CC, et al. (2022) OMiCC: An expanded and enhanced platform for meta-analysis of public gene expression data. STAR protocols, 3(3), 101474.

Shah N, et al. (2016) A crowdsourcing approach for reusing and meta-analyzing gene expression data. Nature biotechnology, 34(8), 803.