

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

EMMA2

RRID:SCR_010940

Type: Tool

Proper Citation

EMMA2 (RRID:SCR_010940)

Resource Information

URL: <http://www.cebitec.uni-bielefeld.de/comics/index.php/emma>

Proper Citation: EMMA2 (RRID:SCR_010940)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented May 17, 2017. A MAGE-compliant software platform for the collaborative analysis and integration of microarray data.

Abbreviations: EMMA2

Synonyms: EMMA 2

Resource Type: service resource

Defining Citation: [PMID:19200358](#)

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: EMMA2

Resource ID: SCR_010940

Alternate IDs: OMICS_00753

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260725T190707+0000

Ratings and Alerts

No rating or validation information has been found for EMMA2.

No alerts have been found for EMMA2.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Droste J, et al. (2020) A maltose-regulated large genomic region is activated by the transcriptional regulator MalT in *Actinoplanes* sp. SE50/110. *Applied microbiology and biotechnology*, 104(21), 9283.

Wolf T, et al. (2017) The MalR type regulator AcrC is a transcriptional repressor of acarbose biosynthetic genes in *Actinoplanes* sp. SE50/110. *BMC genomics*, 18(1), 562.

Rieke S, et al. (2017) Mixture effects of azole fungicides on the adrenal gland in a broad dose range. *Toxicology*, 385, 28.

Toepel J, et al. (2011) Construction and evaluation of a whole genome microarray of *Chlamydomonas reinhardtii*. *BMC genomics*, 12, 579.

Schröder J, et al. (2010) The Zur regulon of *Corynebacterium glutamicum* ATCC 13032. *BMC genomics*, 11, 12.

Rückert C, et al. (2008) The dual transcriptional regulator CysR in *Corynebacterium glutamicum* ATCC 13032 controls a subset of genes of the McbR regulon in response to the availability of sulphide acceptor molecules. *BMC genomics*, 9, 483.