

# Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

## EMBRYS

RRID:SCR\_006689

Type: Tool

### Proper Citation

EMBRYS (RRID:SCR\_006689)

### Resource Information

**URL:** <https://www.embrys.jp/embrys/html/About.html>

**Proper Citation:** EMBRYS (RRID:SCR\_006689)

**Description:** Data collection of gene expression patterns mapped in whole-mount mouse embryo (ICR strain) of mid-gestational stages (Embryonic Day 9.5, 10.5, 11.5), in which most striking dynamics in pattern formation and organogenesis is observed. Collection of gene expression patterns of transcription factors (TFs) and TF-related factors such as transcription cofactors. Genes were extracted from databases including RIKEN Transcription Factor Database and Panther Classification System.

**Abbreviations:** EMBRYS

**Synonyms:** Embryonic Gene Expression Database for Biomedical Research Source, Embryonic gene expression Database as a Biomedical Research Source

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

**Keywords:** Gene, expression, pattern, mapped, whole mount, mouse, embryo, ICR strain, mid gestational stage, transcription, factor, cofactor, data

**Funding:** Japanese Ministry of Education Culture Sports Science and Technology MEXT ; Japanese Ministry of Health Labor and Welfare

**Availability:** Free, Freely available

**Resource Name:** EMBRYS

**Resource ID:** SCR\_006689

**Alternate IDs:** nlx\_153839

**Old URLs:** <http://embrys.jp/embrys/html/MainMenu.html>

**Record Creation Time:** 20220129T080237+0000

**Record Last Update:** 20260912T200143+0000

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## Ratings and Alerts

No rating or validation information has been found for EMBRYS.

No alerts have been found for EMBRYS.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 8 mentions in open access literature.

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

Gamart J, et al. (2021) SMAD4 target genes are part of a transcriptional network that integrates the response to BMP and SHH signaling during early limb bud patterning. Development (Cambridge, England), 148(23).

Yokoyama S, et al. (2017) Analysis of transcription factors expressed at the anterior mouse limb bud. PloS one, 12(5), e0175673.

Motohashi T, et al. (2016) Gene array analysis of neural crest cells identifies transcription factors necessary for direct conversion of embryonic fibroblasts into neural crest cells. Biology open, 5(3), 311.

Bhat R, et al. (2016) Deep phylogenomics of a tandem-repeat galectin regulating appendicular skeletal pattern formation. BMC evolutionary biology, 16(1), 162.

Clarkson MD, et al. (2016) Representation of anatomy in online atlases and databases: a survey and collection of patterns for interface design. BMC developmental biology, 16(1), 18.

Ogura Y, et al. (2015) Identification of HOXD4 Mutations in Spinal Extradural Arachnoid Cyst. PloS one, 10(11), e0142126.

Shimizu H, et al. (2013) The AERO system: a 3D-like approach for recording gene expression patterns in the whole mouse embryo. PloS one, 8(10), e75754.

Ito Y, et al. (2012) A systems approach and skeletal myogenesis. Comparative and functional genomics, 2012, 759407.