

Resource Summary Report

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

miROrtho: the catalogue of animal microRNA genes

RRID:SCR_007797

Type: Tool

Proper Citation

miROrtho: the catalogue of animal microRNA genes (RRID:SCR_007797)

Resource Information

URL: <http://cegg.unige.ch/mirortho>

Proper Citation: miROrtho: the catalogue of animal microRNA genes (RRID:SCR_007797)

Description: It contains predictions of precursor miRNA genes covering several animal genomes combining orthology and a Support Vector Machine. We provide homology extended alignments of already known miRBase families and putative miRNA families exclusively predicted by our SVM and orthology pipeline. The current release of miROrtho covers 46 animal genomes. We provide homology extended alignments of already known miRBase families and putative miRNA families exclusively predicted by our SVM and orthology pipeline.

Synonyms: miROrtho

Resource Type: database, data or information resource

Keywords: bio.tools

Resource Name: miROrtho: the catalogue of animal microRNA genes

Resource ID: SCR_007797

Alternate IDs: nif-0000-03139, biotools:mirortho

Alternate URLs: <https://bio.tools/mirortho>

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260725T191149+0000

Ratings and Alerts

No rating or validation information has been found for miROrtho: the catalogue of animal microRNA genes.

No alerts have been found for miROrtho: the catalogue of animal microRNA genes.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Duvaud S, et al. (2021) Expasy, the Swiss Bioinformatics Resource Portal, as designed by its users. Nucleic acids research, 49(W1), W216.

Hernández-Romero IA, et al. (2019) The Regulatory Roles of Non-coding RNAs in Angiogenesis and Neovascularization From an Epigenetic Perspective. Frontiers in oncology, 9, 1091.

Mishra S, et al. (2015) Circulating miRNAs revealed as surrogate molecular signatures for the early detection of breast cancer. Cancer letters, 369(1), 67.

, et al. (2014) The common marmoset genome provides insight into primate biology and evolution. Nature genetics, 46(8), 850.