

Resource Summary Report

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MIRIAM: Minimal Information Required In the Annotation of Models

RRID:SCR_010547

Type: Tool

Proper Citation

MIRIAM: Minimal Information Required In the Annotation of Models (RRID:SCR_010547)

Resource Information

URL: <http://biomodels.net/miriam/>

Proper Citation: MIRIAM: Minimal Information Required In the Annotation of Models (RRID:SCR_010547)

Description: MIRIAM is an effort to standardise the Minimal Information Required In the Annotation of Models, so that different groups can collaborate on annotating and curating computational models in biology. The goal of the project, initiated by the BioModels.net effort is to produce a set of guidelines suitable for use with any structured format for computational models. MIRIAM is a registered project of the MIBBI (Minimum Information for Biological and Biomedical Investigations). If you are looking for the online resources providing support to MIRIAM Standard, please go to: MIRIAM Resources.

Abbreviations: MIRIAM

Resource Type: data or information resource, topical portal, portal

Resource Name: MIRIAM: Minimal Information Required In the Annotation of Models

Resource ID: SCR_010547

Alternate IDs: nlx_31750

Record Creation Time: 20220129T080259+0000

Record Last Update: 20260811T043401+0000

Ratings and Alerts

No rating or validation information has been found for MIRIAM: Minimal Information Required In the Annotation of Models.

No alerts have been found for MIRIAM: Minimal Information Required In the Annotation of Models.

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](#) .

McKinley ET, et al. (2022) MIRIAM: A machine and deep learning single-cell segmentation and quantification pipeline for multi-dimensional tissue images. Cytometry. Part A : the journal of the International Society for Analytical Cytology, 101(6), 521.

Dreher F, et al. (2012) DIPSBBC--data integration platform for systems biology collaborations. BMC bioinformatics, 13, 85.