

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

Multiple Myeloma survival predictor

RRID:SCR_017651

Type: Tool

Proper Citation

Multiple Myeloma survival predictor (RRID:SCR_017651)

Resource Information

URL: <https://www.synapse.org/#!/Synapse:syn11459638>

Proper Citation: Multiple Myeloma survival predictor (RRID:SCR_017651)

Description: Dockerized environment for winning algorithm in 2017 Multiple Myeloma DREAM Challenge, Sub-Challenge 3.

Resource Type: software resource, data access protocol, web service, service resource

Keywords: Multiple Myeloma, prognostic model, survival analysis, GuanRank, bio.tools

Related Condition: Multiple Myeloma

Availability: Restricted

Resource Name: Multiple Myeloma survival predictor

Resource ID: SCR_017651

Alternate IDs: biotools:Multiple_Myeloma_survival_prediction

Alternate URLs: https://bio.tools/Multiple_Myeloma_survival_prediction

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260808T190109+0000

Ratings and Alerts

No rating or validation information has been found for Multiple Myeloma survival predictor.

No alerts have been found for Multiple Myeloma survival predictor.

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 [SPARC SAWG](#) .

Tomasoni M, et al. (2020) MONET: a toolbox integrating top-performing methods for network modularization. Bioinformatics (Oxford, England), 36(12), 3920.

Sun C, et al. (2019) Prognostic model for multiple myeloma progression integrating gene expression and clinical features. GigaScience, 8(12).