

Resource Summary Report

Generated by SPARC SAWG on Aug 13, 2026

Multiple Myeloma Genomics Portal

RRID:SCR_003722

Type: Tool

Proper Citation

Multiple Myeloma Genomics Portal (RRID:SCR_003722)

Resource Information

URL: <http://www.broadinstitute.org/mmgp/>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on January 6, 2023. Database providing access and limited analysis to the MMGP portal data sets. These include the MMRC funded reference array comparative genomic hybridization (aCGH) and gene expression data and additional public multiple myeloma datasets. The MMGP will be updated with new features such as additional data and analysis tools as they become available.

Abbreviations: MMGP

Synonyms: MMRC Multiple Myeloma Genomics Portal

Resource Type: data or information resource, database

Keywords: gene, genomics, mutation, single-nucleotide polymorphism, array comparative genomic hybridization, gene expression, resequencing, rai, sample annotation, differential expression

Related Condition: Multiple myeloma

Funding: Multiple Myeloma Research Foundation

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Multiple Myeloma Genomics Portal

Resource ID: SCR_003722

Alternate IDs: nlx_157900

Record Creation Time: 20220129T080220+0000

Ratings and Alerts

No rating or validation information has been found for Multiple Myeloma Genomics Portal.

No alerts have been found for Multiple Myeloma Genomics Portal.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Qu K, et al. (2016) Integrative genomic analysis by interoperation of bioinformatics tools in GenomeSpace. Nature methods, 13(3), 245.

Cottini F, et al. (2014) Rescue of Hippo coactivator YAP1 triggers DNA damage-induced apoptosis in hematological cancers. Nature medicine, 20(6), 599.

Delmore JE, et al. (2011) BET bromodomain inhibition as a therapeutic strategy to target c-Myc. Cell, 146(6), 904.