

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

MethMarker

RRID:SCR_010908

Type: Tool

Proper Citation

MethMarker (RRID:SCR_010908)

Resource Information

URL: <http://methmarker.mpi-inf.mpg.de/>

Proper Citation: MethMarker (RRID:SCR_010908)

Description: Tool that facilitates the design and optimization of gene-specific DNA methylation assays. Beyond its use as an epigenetic primer-design tool, it provides extensive support for epigenetic biomarker optimization. Download MethMarker or start it directly from within your web browser.

Abbreviations: MethMarker

Resource Type: software resource

Defining Citation: [PMID:19804638](#)

Keywords: dna methylation, bio.tools

Availability: Acknowledgement requested

Resource Name: MethMarker

Resource ID: SCR_010908

Alternate IDs: OMICS_00636, biotools:methmarker

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

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260919T195202+0000

Ratings and Alerts

No rating or validation information has been found for MethMarker.

No alerts have been found for MethMarker.

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 [nidm-terms](#) .

Sandoval AQ, et al. (2025) Nuclear adenomatous polyposis coli elevates STAT1 and reduces CXCL1,2, and 3 expression and inhibits neutrophil recruitment. Cellular signalling, 134, 111957.

Dooley SA, et al. (2025) The Molecular Motor Myosin 5B and Its Folding Chaperone UNC45A Are Decreased in Colorectal Cancer. Cellular and molecular gastroenterology and hepatology, 19(9), 101537.

Schiffman P, et al. (2009) MethMarker: user-friendly design and optimization of gene-specific DNA methylation assays. Genome biology, 10(10), R105.