

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org) on Apr 9, 2025

BatMeth

RRID:SCR_010906

Type: Tool

Proper Citation

BatMeth (RRID:SCR_010906)

Resource Information

URL: <https://code.google.com/p/batmeth/>

Proper Citation: BatMeth (RRID:SCR_010906)

Description: Improved mapper for bisulfite sequencing reads on DNA methylation.

Abbreviations: BatMeth

Resource Type: software resource

Keywords: bio.tools

Funding:

Resource Name: BatMeth

Resource ID: SCR_010906

Alternate IDs: biotools:batmeth, OMICS_00574

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

Record Creation Time: 20220129T080301+0000

Record Last Update: 20250214T183147+0000

Ratings and Alerts

No rating or validation information has been found for BatMeth.

No alerts have been found for BatMeth.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Liu Y, et al. (2019) H3K4me2 functions as a repressive epigenetic mark in plants. *Epigenetics & chromatin*, 12(1), 40.