

Resource Summary Report

Generated by [nidm-terms](#) on Aug 9, 2026

CandiMeth

RRID:SCR_017974

Type: Tool

Proper Citation

CandiMeth (RRID:SCR_017974)

Resource Information

URL: <http://bit.do/canidmeth-github>

Proper Citation: CandiMeth (RRID:SCR_017974)

Description: Software tool for visualisation and quantification of DNA methylation at candidate features.

Abbreviations: CandiMeth

Synonyms: CANDIdate feature METHylation

Resource Type: data analysis software, service resource, data processing software, software application, analysis service resource, software resource, production service resource

Keywords: DNA, methylation, candidate feature, visualisation, quantification, bio.tools

Funding: Medical Research Council ;
ESRC/BBSRC

Availability: Free, Freely available

Resource Name: CandiMeth

Resource ID: SCR_017974

Alternate IDs: biotools:CandiMeth

Alternate URLs: <http://bit.do/candimeth>, <https://usegalaxy.org/u/sarajayne-thursby/w/candimeth-8>, <https://bio.tools/CandiMeth>

License: GNU General Public License v3.0

Record Creation Time: 20220129T080338+0000

Ratings and Alerts

No rating or validation information has been found for CandiMeth.

No alerts have been found for CandiMeth.

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 [nidm-terms](#) .

Hsieh ML, et al. (2025) Elucidating the Mechanism of Radiation Therapy on Mesenchymal Cell Fate in Preventing Heterotopic Ossification. Annals of surgery, 282(6), 1140.

Thursby SJ, et al. (2020) CandiMeth: Powerful yet simple visualization and quantification of DNA methylation at candidate genes. GigaScience, 9(6).