Resource Summary Report

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

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

Record Last Update: 20250409T061540+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 1 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

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