Resource Summary Report

Generated by NIF on May 2, 2025

MethylCoder

RRID:SCR_012997

Type: Tool

Proper Citation

MethylCoder (RRID:SCR_012997)

Resource Information

URL: https://github.com/brentp/methylcode

Proper Citation: MethylCoder (RRID:SCR_012997)

Description: A single program that takes of bisulfite-treated reads and outputs per-base

methylation data.

Abbreviations: MethylCoder

Resource Type: software resource

Defining Citation: PMID:21724594

Keywords: bio.tools

Funding:

Resource Name: MethylCoder

Resource ID: SCR_012997

Alternate IDs: biotools:methylcoder, OMICS_00585

Alternate URLs: https://bio.tools/methylcoder

Record Creation Time: 20220129T080313+0000

Record Last Update: 20250420T014628+0000

Ratings and Alerts

No rating or validation information has been found for MethylCoder.

No alerts have been found for MethylCoder.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Han Y, et al. (2016) Integrating Epigenomics into the Understanding of Biomedical Insight. Bioinformatics and biology insights, 10, 267.

Baheti S, et al. (2016) Targeted alignment and end repair elimination increase alignment and methylation measure accuracy for reduced representation bisulfite sequencing data. BMC genomics, 17, 149.

Karcher JR, et al. (2015) Genome-wide epigenetic and proteomic analysis reveals altered Notch signaling in EPC dysfunction. Physiological reports, 3(4).

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database: the journal of biological databases and curation, 2015.