Resource Summary Report

Generated by RRID on Apr 28, 2025

MethBase

RRID:SCR_017487
Type: Tool

Proper Citation

MethBase (RRID:SCR_017487)

Resource Information

URL: http://smithlabresearch.org/software/methbase/

Proper Citation: MethBase (RRID:SCR_017487)

Description: Central reference methylome database created from public BS-seq datasets. Provides methylation level at individual sites, regions of allele specific methylation, hypo- or hyper-methylated regions, partially methylated regions, and detailed meta data and summary statistics.

Synonyms: MethBase: a reference methylome database

Resource Type: service resource, database, data or information resource

Keywords: Methylome, database, public, BSseq, dataset, methylation, site, region, allele, specific, metadata, statistics, bio.tools

Funding:

Availability: Free, Freely available

Resource Name: MethBase

Resource ID: SCR_017487

Alternate IDs: BioTools:MethBase, biotools:Methbase

Alternate URLs: https://bio.tools/MethBase, https://bio.tools/MethBase, https://bio.tools/MethBase

Record Creation Time: 20220129T080335+0000

Record Last Update: 20250428T054053+0000

Ratings and Alerts

No rating or validation information has been found for MethBase.

No alerts have been found for MethBase.

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 RRID.

Ehrlich M, et al. (2024) Epigenetics of Genes Preferentially Expressed in Dissimilar Cell Populations: Myoblasts and Cerebellum. Epigenomes, 8(1).