Resource Summary Report

Generated by dkNET on Aug 17, 2026

CanMETH database

RRID:SCR_023947
Type: Tool

Proper Citation

CanMETH database (RRID:SCR_023947)

Resource Information

URL: http://meth.liclab.net/CanMethdb/

Proper Citation: CanMETH database (RRID:SCR_023947)

Description: Comprehensive DNA methylation database. Provides genome wide mapping of methylation sites to target genes through integrating a large number of DNA methylation and gene expression profiles in human diseases, and to annotate methylation sites to abundant regulatory elements and TFs.

Resource Type: data or information resource, portal, database, project portal

Keywords: Hengyang Medical School, University of South China Hengyang, DNA methylation data, DNA methylation, gene expression profiles, human diseases, annotate methylation sites, abundant regulatory elements,

Availability: Free, Freely available

Resource Name: CanMETH database

Resource ID: SCR_023947

Record Creation Time: 20230818T050222+0000

Record Last Update: 20260817T040704+0000

Ratings and Alerts

No rating or validation information has been found for CanMETH database .

No alerts have been found for CanMETH database .

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 dkNET .

Bisht D, et al. (2026) Liquid biopsy-based diagnostic evaluation of hypermethylated CpG sites for ovarian cancer diagnosis. Molecular oncology.