Resource Summary Report

Generated by NIF on May 21, 2025

MetMap

RRID:SCR_006954
Type: Tool

Proper Citation

MetMap (RRID:SCR_006954)

Resource Information

URL: http://math.mcb.berkeley.edu/~meromit/MetMap/

Proper Citation: MetMap (RRID:SCR_006954)

Description: A computational pipeline for the analysis of MethylSeq experiments.

Abbreviations: MetMap

Resource Type: software resource

Funding:

Resource Name: MetMap

Resource ID: SCR_006954

Alternate IDs: OMICS_00618

Record Creation Time: 20220129T080239+0000

Record Last Update: 20250519T203453+0000

Ratings and Alerts

No rating or validation information has been found for MetMap.

No alerts have been found for MetMap.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Dimitrieva S, et al. (2025) Biologically relevant integration of transcriptomics profiles from cancer cell lines, patient-derived xenografts, and clinical tumors using deep learning. Science advances, 11(3), eadn5596.

Guo Y, et al. (2024) IGSF3 is a homophilic cell adhesion molecule that drives lung metastasis of melanoma by promoting adhesion to vascular endothelium. Cancer science, 115(6), 1936.

Schneider C, et al. (2024) A Novel AMPK Inhibitor Sensitizes Pancreatic Cancer Cells to Ferroptosis Induction. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 11(31), e2307695.

Gan S, et al. (2023) Distinct tumor architectures for metastatic colonization of the brain. bioRxiv : the preprint server for biology.

Mathur D, et al. (2023) The Ratio of Key Metabolic Transcripts Is a Predictive Biomarker of Breast Cancer Metastasis to the Lung. Cancer research, 83(20), 3478.

Zhu Q, et al. (2015) EHR based Genetic Testing Knowledge Base (iGTKB) Development. BMC medical informatics and decision making, 15 Suppl 4(Suppl 4), S3.