

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

[NGSmethPipe](#)

RRID:SCR_000583

Type: Tool

Proper Citation

NGSmethPipe (RRID:SCR_000583)

Resource Information

URL: <http://bioinfo2.ugr.es/NGSmethPipe/>

Proper Citation: NGSmethPipe (RRID:SCR_000583)

Description: A software tool which generates high-quality methylation maps.

Abbreviations: NGSmethPipe

Synonyms: NGSmethPipe - A tool to generate high-quality methylation maps

Resource Type: software resource

Keywords: computation, genomics, bioinformatics, methylation maps, visualization

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: NGSmethPipe

Resource ID: SCR_000583

Alternate IDs: OMICS_00611

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260801T190115+0000

Ratings and Alerts

No rating or validation information has been found for NGSmethPipe.

No alerts have been found for NGSmethPipe.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

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

Geisen S, et al. (2014) NGSmethDB: an updated genome resource for high quality, single-cytosine resolution methylomes. Nucleic acids research, 42(Database issue), D53.

Barturen G, et al. (2013) CpGislandEVO: a database and genome browser for comparative evolutionary genomics of CpG islands. BioMed research international, 2013, 709042.