

# Resource Summary Report

Generated by T1D on Aug 3, 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](#)

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [T1D](#) .

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.