

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

DamMet

RRID:SCR_016959

Type: Tool

Proper Citation

DamMet (RRID:SCR_016959)

Resource Information

URL: <https://gitlab.com/KHanghoj/DamMet>

Proper Citation: DamMet (RRID:SCR_016959)

Description: Software tool as a full probabilistic model for mapping ancient methylomes using sequencing data underlying an ancient specimen.

Resource Type: software resource, software application, data processing software, service resource

Keywords: mapping, sequencing, data, ancient, specimen, DNA, methylation

Availability: Free, Freely available, Tutorial available

Resource Name: DamMet

Resource ID: SCR_016959

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Record Creation Time: 20220129T080332+0000

Record Last Update: 20260808T190321+0000

Ratings and Alerts

No rating or validation information has been found for DamMet.

No alerts have been found for DamMet.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Seguin-Orlando A, et al. (2021) Heterogeneous Hunter-Gatherer and Steppe-Related Ancestries in Late Neolithic and Bell Beaker Genomes from Present-Day France. *Current biology : CB*, 31(5), 1072.

Hanghøj K, et al. (2019) DamMet: ancient methylome mapping accounting for errors, true variants, and post-mortem DNA damage. *GigaScience*, 8(4).