mapDamage
RRID:SCR_001240
Type: Tool

Proper Citation
mapDamage (RRID:SCR_001240)

Resource Information

URL: http://ginolhac.github.io/mapDamage/

Description: Software for tracking and quantifying DNA damage patterns among ancient DNA sequencing reads generated by Next-Generation Sequencing platforms.

Resource Name: mapDamage

Proper Citation: mapDamage (RRID:SCR_001240)

Resource Type: Resource, software resource

Keywords: python, r, illumina, windows, perl, dna damage, dna sequencing, next-generation sequencing, dna

Resource ID: SCR_001240

Parent Organization: University of Copenhagen; Copenhagen; Denmark

References: PMID: 23613487, PMID: 21659319

Website Status: Last checked up

Alternate IDs: OMICS_02099

Abbreviations: mapDamage

Mentions Count: 99

Ratings and Alerts
No rating or validation information has been found for mapDamage.

No alerts have been found for mapDamage.

Data and Source Information
Source: SciCrunch Registry

Usage and Citation Metrics

We found 99 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](#).

Csáky V, et al. (2020) Genetic insights into the social organisation of the Avar period elite in the 7th century AD Carpathian Basin. Scientific reports, 10(1), 948.


Wales N, et al. (2019) Ancient DNA reveals the timing and persistence of organellar genetic bottlenecks over 3,000 years of sunflower domestication and improvement. Evolutionary
applications, 12(1), 38-53.


