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: 95

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

No alerts have been found for mapDamage.

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**Data and Source Information**

**Source:** SciCrunch Registry

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**Usage and Citation Metrics**

We found 95 mentions in open access literature.

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


Velsko IM, et al. (2019) Microbial differences between dental plaque and historic dental calculus are related to oral biofilm maturation stage. Microbiome, 7(1), 102.


