PALEOMIX
RRID:SCR_015057
Type: Tool

Proper Citation

PALEOMIX (RRID:SCR_015057)

Resource Information

URL: https://github.com/MikkelSchubert/paleomix

Proper Citation: PALEOMIX (RRID:SCR_015057)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on February 28, 2023. Software toolkit for the processing of ancient and modern HTS data. PALEOMIX also aids in metagenomic analysis of the extracts from the HTS processing.

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

Defining Citation: PMID:24722405, DOI:10.1038/nprot.2014.063

Keywords: hts data, high-throughput sequencing, ancient dna, adna, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: PALEOMIX

Resource ID: SCR_015057

Alternate IDs: biotools:paleomix, OMICS_03749


Record Creation Time: 20220129T080323+0000

Record Last Update: 20240702T054002+0000

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

No alerts have been found for PALEOMIX.

**Data and Source Information**

**Source:** [SciCrunch Registry](SciCrunch.org)

**Usage and Citation Metrics**

We found 55 mentions in open access literature.

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

Gavriil M, et al. (2024) 2-Hydroxyglutarate modulates histone methylation at specific loci and alters gene expression via Rph1 inhibition. Life science alliance, 7(2).


Breglia F, et al. (2023) Disentangling the origins of viticulture in the western Mediterranean. Scientific reports, 13(1), 17284.


Slimak L, et al. (2022) Modern human incursion into Neanderthal territories 54,000 years ago at Mandrin, France. Science advances, 8(6), eabj9496.


Canales NA, et al. (2022) A highly contiguous, scaffold-level nuclear genome assembly for the fever tree (Cinchona pubescens Vahl) as a novel resource for Rubiaceae research. GigaByte (Hong Kong, China), 2022, gigabyte71.
