

Resource Summary Report

Generated by SPARC SAWG on Sep 19, 2026

DELIMINATE

RRID:SCR_008956

Type: Tool

Proper Citation

DELIMINATE (RRID:SCR_008956)

Resource Information

URL: <http://metagenomics.atc.tcs.com/compression/DELIMINATE/>

Proper Citation: DELIMINATE (RRID:SCR_008956)

Description: A practical implementation of a novel compression approach that can rapidly compress FASTA files containing genomic sequence data in a loss-less fashion.

Abbreviations: DELIMINATE

Synonyms: DELIMINATE: A fast and efficient method for loss-less compression of genomic sequences

Resource Type: software resource

Defining Citation: [PMID:22833526](#)

Resource Name: DELIMINATE

Resource ID: SCR_008956

Alternate IDs: OMICS_00953

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260919T195147+0000

Ratings and Alerts

No rating or validation information has been found for DELIMINATE.

No alerts have been found for DELIMINATE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Kryukov K, et al. (2019) Nucleotide Archival Format (NAF) enables efficient lossless reference-free compression of DNA sequences. Bioinformatics (Oxford, England), 35(19), 3826.