

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 12, 2026

DEXTRACTOR

RRID:SCR_024005

Type: Tool

Proper Citation

DEXTRACTOR (RRID:SCR_024005)

Resource Information

URL: <https://github.com/thegenemyers/DEXTRACTOR>

Proper Citation: DEXTRACTOR (RRID:SCR_024005)

Description: Software as Bax file decoder and data compressor.

Synonyms: Dextractor and Compression Command Library, dextractor

Resource Type: software resource, source code

Keywords: Bax, file decoder, data compressor,

Availability: Free, Available for download, Freely available

Resource Name: DEXTRACTOR

Resource ID: SCR_024005

Alternate URLs: <https://sources.debian.org/src/dextractor/>

Record Creation Time: 20230824T050211+0000

Record Last Update: 20260912T200428+0000

Ratings and Alerts

No rating or validation information has been found for DEXTRACTOR.

No alerts have been found for DEXTRACTOR.

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](#) .

Chetruengchai W, et al. (2024) De novo genome assembly and transcriptome sequencing in foot and mantle tissues of *Megaustenia siamensis* reveals components of adhesive substances. *Scientific reports*, 14(1), 13756.

Korhonen PK, et al. (2019) Common workflow language (CWL)-based software pipeline for de novo genome assembly from long- and short-read data. *GigaScience*, 8(4).