

Resource Summary Report

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Cdbfasta

RRID:SCR_016057

Type: Tool

Proper Citation

Cdbfasta (RRID:SCR_016057)

Resource Information

URL: <http://cdbfasta.sourceforge.net/>

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Description: Software tool for indexing and retrieval of nucleotide sequences from FASTA (DNA and protein sequence alignment software) record databases. It has the option to compress data records.

Abbreviations: CDB

Synonyms: Constant DataBase

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

Keywords: index, retrieval, nucleotide, sequence, fasta, database, multi file, compress, record

Funding: The Free Software Foundation (FSF)

Availability: Free, Available for download

Resource Name: Cdbfasta

Resource ID: SCR_016057

Alternate IDs: OMICS_19793

Alternate URLs: <https://github.com/gperte/cdbfasta>,
<https://sources.debian.org/src/cdbfasta/>

License: Artistic License 2.0

Record Creation Time: 20220129T080328+0000

Record Last Update: 20260731T042958+0000

Ratings and Alerts

No rating or validation information has been found for Cdbfasta.

No alerts have been found for Cdbfasta.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.