

# Resource Summary Report

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## Cdbfasta

RRID:SCR\_016057

Type: Tool

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### Proper Citation

Cdbfasta (RRID:SCR\_016057)

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### 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:** software application, software toolkit, data transfer software, data processing software, data compression software, software resource

**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/gpertea/cdbfasta>,  
<https://sources.debian.org/src/cdbfasta/>

**License:** Artistic License 2.0

**Record Creation Time:** 20220129T080328+0000

**Record Last Update:** 20250519T203921+0000

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## Ratings and Alerts

No rating or validation information has been found for Cdbfasta.

No alerts have been found for Cdbfasta.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.