

Resource Summary Report

Generated by [PRECISE-TBI](#) on Jul 31, 2026

PREFAB

RRID:SCR_001009

Type: Tool

Proper Citation

PREFAB (RRID:SCR_001009)

Resource Information

URL: <http://www.drive5.com/muscle/prefab.htm>

Proper Citation: PREFAB (RRID:SCR_001009)

Description: Downloadable data designed for testing multiple sequence alignment methods.

Abbreviations: PREFAB

Synonyms: PREFAB - Protein Reference Alignment Benchmark, Protein Reference Alignment Benchmark

Resource Type: data or information resource, database

Defining Citation: [PMID:15034147](#)

Keywords: database, downloadable data, sequence alignment, sequence alignment method

Availability: Free, Available for download, Freely available

Resource Name: PREFAB

Resource ID: SCR_001009

Alternate IDs: OMICS_00984

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260729T044006+0000

Ratings and Alerts

No rating or validation information has been found for PREFAB.

No alerts have been found for PREFAB.

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 [PRECISE-TBI](#) .

Pei J, et al. (2006) MUMMALS: multiple sequence alignment improved by using hidden Markov models with local structural information. Nucleic acids research, 34(16), 4364.