

# Resource Summary Report

Generated by SPARC SAWG on Sep 16, 2026

## PREFAB

RRID:SCR\_001009

Type: Tool

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### Proper Citation

PREFAB (RRID:SCR\_001009)

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### 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:** 20260912T200121+0000

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### Ratings and Alerts

No rating or validation information has been found for PREFAB.

No alerts have been found for PREFAB.

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## Data and Source Information

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

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## 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](#) .

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.