

# Resource Summary Report

Generated by [NIF](#) on Aug 9, 2026

## Protein Research Foundation

RRID:SCR\_006871

Type: Tool

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### Proper Citation

Protein Research Foundation (RRID:SCR\_006871)

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### Resource Information

**URL:** <http://www.prf.or.jp/index-e.html>

**Proper Citation:** Protein Research Foundation (RRID:SCR\_006871)

**Description:** The Protein Research Foundation has collected information related to amino acids, peptides and proteins and created several databases. The website is partially commercial and some databases are available only for a fee, although all can be accessed for a brief period as a guest account as well. They provide a literature search database, a Peptide/Protein Sequence Database, and a Synthetic Compounds Database.

**Synonyms:** PRF

**Resource Type:** database, data or information resource, portal, topical portal

**Resource Name:** Protein Research Foundation

**Resource ID:** SCR\_006871

**Alternate IDs:** nif-0000-03335

**Old URLs:** <http://www.prf.or.jp/en/index.shtml>

**Record Creation Time:** 20220129T080238+0000

**Record Last Update:** 20260808T185903+0000

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### Ratings and Alerts

No rating or validation information has been found for Protein Research Foundation.

No alerts have been found for Protein Research Foundation.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [NIF](#) .

Mora-Márquez F, et al. (2025) gymnotoa-db: a database and application to optimize functional annotation in gymnosperms. Database : the journal of biological databases and curation, 2025.

Chen J, et al. (2014) Transcriptome sequencing and profiling of expressed genes in phloem and xylem of ramie (*Boehmeria nivea* L. Gaud). PloS one, 9(10), e110623.

Mariño-Ramírez L, et al. (2011) The Histone Database: an integrated resource for histones and histone fold-containing proteins. Database : the journal of biological databases and curation, 2011, bar048.