

# Resource Summary Report

Generated by PRECISE-TBI on Sep 15, 2026

## KnowPredsite

RRID:SCR\_011974

Type: Tool

### Proper Citation

KnowPredsite (RRID:SCR\_011974)

### Resource Information

**URL:** <http://bio-cluster.iis.sinica.edu.tw/kbloc/index.html>

**Proper Citation:** KnowPredsite (RRID:SCR\_011974)

**Description:** A knowledge-based data analysis service to predict the localization site(s) of both single-localized and multi-localized proteins.

**Abbreviations:** KnowPredsite

**Synonyms:** KnowPredsite: A web server for predicting single and multiple subcellular localization sites, KnowPred site, KnowPredsite: A general protein subcellular localization predictor for eukaryotes and prokaryotes

**Resource Type:** analysis service resource, data analysis service, production service resource, service resource

**Defining Citation:** [PMID:19958518](#)

**Keywords:** homolog, protein, subcellular localization

**Resource Name:** KnowPredsite

**Resource ID:** SCR\_011974

**Alternate IDs:** OMICS\_01625

**Record Creation Time:** 20220129T080307+0000

**Record Last Update:** 20260912T200207+0000

### Ratings and Alerts

No rating or validation information has been found for KnowPredsite.

No alerts have been found for KnowPredsite.

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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 [PRECISE-TBI](#) .

Gupta SK, et al. (2021) An Ebola, Neisseria and Trypanosoma human protein interaction census reveals a conserved human protein cluster targeted by various human pathogens. Computational and structural biotechnology journal, 19, 5292.