

Resource Summary Report

Generated by T1D on Aug 8, 2026

Proteome Analyst PA-GOSUB

RRID:SCR_008234

Type: Tool

Proper Citation

Proteome Analyst PA-GOSUB (RRID:SCR_008234)

Resource Information

URL: <http://www.cs.ualberta.ca/~bioinfo/PA/GOSUB/>

Proper Citation: Proteome Analyst PA-GOSUB (RRID:SCR_008234)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented on June 30, 2015. Refer to Proteome Analyst 3.0. Subcellular Localization and GO General Molecular Function predictions for many model organism proteomes using Protein Analyst, with a very high coverage rate. When users blast their proteins against the database of results, they will not only be shown blast homologs from the model organisms, but also the Subcellular Localization and GO General Molecular Function predictions as well.

Abbreviations: PA-GOSUB

Synonyms: Proteome Analyst

Resource Type: data or information resource, software resource, data set

Keywords: subcellular localization, homolog, localization, molecular, organism, prediction, protein, proteome, subcellular, organelle, gene ontology cellular component, nucleus, golgi apparatus, mitochondrion

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Proteome Analyst PA-GOSUB

Resource ID: SCR_008234

Alternate IDs: nif-0000-21342

Record Creation Time: 20220129T080246+0000

Record Last Update: 20260808T190313+0000

Ratings and Alerts

No rating or validation information has been found for Proteome Analyst PA-GOSUB.

No alerts have been found for Proteome Analyst PA-GOSUB.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

González-Fernández R, et al. (2010) Proteomics of plant pathogenic fungi. Journal of biomedicine & biotechnology, 2010, 932527.

Coulibaly I, et al. (2008) Bioinformatic tools for inferring functional information from plant microarray data II: Analysis beyond single gene. International journal of plant genomics, 2008, 893941.