

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](#) on May 13, 2025

MetaLocGramN

RRID:SCR_003154

Type: Tool

Proper Citation

MetaLocGramN (RRID:SCR_003154)

Resource Information

URL: <http://iimcb.genesilico.pl/MetaLocGramN/>

Proper Citation: MetaLocGramN (RRID:SCR_003154)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on January 5, 2023. A tool for subcellular localization prediction of Gram-negative proteins. You can also use MetaGramLocN via SOAP. SOAP enables you to invoke our method from scripts written in your programming language of choice.

Abbreviations: MetaLocGramN

Resource Type: web service, software resource, data access protocol, analysis service resource, production service resource, service resource, data analysis service

Defining Citation: [PMID:22705560](#)

Keywords: subcellular localization, protein, prediction, sequence, analysis, gram-negative protein, gram-negative, gram-negative bacteria

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: MetaLocGramN

Resource ID: SCR_003154

Alternate IDs: OMICS_01626

Record Creation Time: 20220129T080217+0000

Record Last Update: 20250513T060507+0000

Ratings and Alerts

No rating or validation information has been found for MetaLocGramN.

No alerts have been found for MetaLocGramN.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Stuart RK, et al. (2016) Light Regimes Shape Utilization of Extracellular Organic C and N in a Cyanobacterial Biofilm. mBio, 7(3).

Rubiano-Labrador C, et al. (2015) Salt Stress Induced Changes in the Exoproteome of the Halotolerant Bacterium Tistlia consotensis Deciphered by Proteogenomics. PloS one, 10(8), e0135065.

Moumène A, et al. (2015) Proteomic profiling of the outer membrane fraction of the obligate intracellular bacterial pathogen Ehrlichia ruminantium. PloS one, 10(2), e0116758.