

# Resource Summary Report

Generated by [dkNET](#) on Aug 26, 2026

## MetaLocGramN

RRID:SCR\_003154

Type: Tool

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### Proper Citation

MetaLocGramN (RRID:SCR\_003154)

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### 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:** analysis service resource, data access protocol, data analysis service, production service resource, service resource, software resource, web service

**Defining Citation:** [PMID:22705560](#)

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

**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:** 20260821T193658+0000

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### Ratings and Alerts

No rating or validation information has been found for MetaLocGramN.

No alerts have been found for MetaLocGramN.

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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 [dkNET](#) .

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

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

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.