

Resource Summary Report

Generated by [Kravitz](#) on Sep 18, 2026

[mGOASVM](#)

RRID:SCR_013098

Type: Tool

Proper Citation

mGOASVM (RRID:SCR_013098)

Resource Information

URL: <http://bioinfo.eie.polyu.edu.hk/mGoaSvmServer/mGOASVM.html>

Proper Citation: mGOASVM (RRID:SCR_013098)

Description: Data analysis service for the prediction of multi-label protein subcellular localization based on gene ontology and support vector machines. Web services are also available.

Abbreviations: mGOASVM

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

Defining Citation: [PMID:23130999](#)

Keywords: subcellular localization, gram-negative protein, virus, protein

Resource Name: mGOASVM

Resource ID: SCR_013098

Alternate IDs: OMICS_01627

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260912T195759+0000

Ratings and Alerts

No rating or validation information has been found for mGOASVM.

No alerts have been found for mGOASVM.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Geng X, et al. (2024) Genome-wide analysis of soybean hypoxia inducible gene domain containing genes: a functional investigation of GmHIGD3. *Frontiers in plant science*, 15, 1403841.

Dutta B, et al. (2021) In silico characterization of bacterial chitinase: illuminating its relationship with archaeal and eukaryotic cousins. *Journal, genetic engineering & biotechnology*, 19(1), 19.

Chakraborty C, et al. (2017) Rising Strengths Hong Kong SAR in Bioinformatics. *Interdisciplinary sciences, computational life sciences*, 9(2), 224.

Wan S, et al. (2012) mGOASVM: Multi-label protein subcellular localization based on gene ontology and support vector machines. *BMC bioinformatics*, 13, 290.