

Resource Summary Report

Generated by [RRID](#) on Apr 8, 2025

PlantLoc

RRID:SCR_003138

Type: Tool

Proper Citation

PlantLoc (RRID:SCR_003138)

Resource Information

URL: <http://cal.tongji.edu.cn/PlantLoc/index.jsp>

Proper Citation: PlantLoc (RRID:SCR_003138)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on January 4, 2023. An accurate web server for predicting plant protein subcellular localization by substantiality motif.

Abbreviations: PlantLoc

Synonyms: PlantLoc: Plant Proteins Subcellular Localization Prediction Server

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

Defining Citation: [PMID:23729470](#)

Keywords: subcellular localization, protein

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: PlantLoc

Resource ID: SCR_003138

Alternate IDs: OMICS_01632

Record Creation Time: 20220129T080217+0000

Record Last Update: 20250407T215341+0000

Ratings and Alerts

No rating or validation information has been found for PlantLoc.

No alerts have been found for PlantLoc.

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 [RRID](#).

Luan H, et al. (2019) Comprehensive Analysis of Soybean Mosaic Virus P3 Protein Interactors and Hypersensitive Response-Like Lesion-Inducing Protein Function. *International journal of molecular sciences*, 20(14).

Figueiredo A, et al. (2017) Specific adjustments in grapevine leaf proteome discriminating resistant and susceptible grapevine genotypes to *Plasmopara viticola*. *Journal of proteomics*, 152, 48.

Chai C, et al. (2015) Soybean transcription factor ORFeome associated with drought resistance: a valuable resource to accelerate research on abiotic stress resistance. *BMC genomics*, 16(1), 596.

Darbani B, et al. (2015) Deciphering Mineral Homeostasis in Barley Seed Transfer Cells at Transcriptional Level. *PloS one*, 10(11), e0141398.