Resource Summary Report

Generated by <u>RRID</u> on Apr 11, 2025

WegoLoc

RRID:SCR_001402 Type: Tool

Proper Citation

WegoLoc (RRID:SCR_001402)

Resource Information

URL: http://www.btool.org/WegoLoc

Proper Citation: WegoLoc (RRID:SCR_001402)

Description: Data analysis service that predicts protein subcellular localizations of animal, fungal, plant, and human proteins based on sequence similarity and gene ontology information.

Abbreviations: WegoLoc

Synonyms: weighted gene ontology term based subcellular locallization prediction

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

Keywords: subcellular localization, protein

Funding:

Resource Name: WegoLoc

Resource ID: SCR_001402

Alternate IDs: OMICS_01636

Record Creation Time: 20220129T080207+0000

Record Last Update: 20250411T054636+0000

Ratings and Alerts

No rating or validation information has been found for WegoLoc.

No alerts have been found for WegoLoc.

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 <u>RRID</u>.

Han Y, et al. (2024) Effects of microgravity on neural crest stem cells. Frontiers in neuroscience, 18, 1379076.

Pernis M, et al. (2023) Secretome analysis revealed that cell wall remodeling and starch catabolism underlie the early stages of somatic embryogenesis in Pinus nigra. Frontiers in plant science, 14, 1225424.

Perutka Z, et al. (2021) Proteome Analysis of Condensed Barley Mitotic Chromosomes. Frontiers in plant science, 12, 723674.

Calvani M, et al. (2018) ?3-Adrenoreceptors Control Mitochondrial Dormancy in Melanoma and Embryonic Stem Cells. Oxidative medicine and cellular longevity, 2018, 6816508.