Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 15, 2025

PEPPER

RRID:SCR_000431
Type: Tool

Proper Citation

PEPPER (RRID:SCR_000431)

Resource Information

URL: http://apps.cytoscape.org/apps/pepper

Proper Citation: PEPPER (RRID:SCR_000431)

Description: A Cytoscape app designed to identify protein pathways / complexes as densely connected subnetworks from seed lists of proteins derived from pull-down assays (i.e AP-MS...).

Abbreviations: PEPPER

Synonyms: Protein complex Expansion using Protein-Protein intERaction networks, Protein complex Expansion using Protein-Protein intERactions

Resource Type: software application, software resource

Defining Citation: PMID:25138169

Keywords: plugin, protein-protein interaction, network, bio.tools

Funding:

Resource Name: PEPPER

Resource ID: SCR_000431

Alternate IDs: biotools:pepper, OMICS_05485

Alternate URLs: https://bio.tools/pepper

Record Creation Time: 20220129T080201+0000

Record Last Update: 20250412T054542+0000

Ratings and Alerts

No rating or validation information has been found for PEPPER.

No alerts have been found for PEPPER.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Wang YS, et al. (2024) Chromosome-level genome assemblies of two littorinid marine snails indicate genetic basis of intertidal adaptation and ancient karyotype evolved from bilaterian ancestors. GigaScience, 13.