

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 3, 2026

PEPPER

RRID:SCR_000431

Type: Tool

Proper Citation

PEPPER (RRID:SCR_000431)

Resource Information

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

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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 resource, software application

Defining Citation: [PMID:25138169](#)

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

Availability: Free, Available for download, Freely available

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: 20260801T191030+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 [PRECISE-TBI](#) .

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.