

Resource Summary Report

Generated by [dkNET](#) on Aug 16, 2026

CoffeeProt

RRID:SCR_022256

Type: Tool

Proper Citation

CoffeeProt (RRID:SCR_022256)

Resource Information

URL: <https://coffeeprot.com/>

Proper Citation: CoffeeProt (RRID:SCR_022256)

Description: Open source software tool for correlation and functional enrichment of proteome wide systems genetics. Analyzes genetic variants associated to protein networks and phenotypic traits. Uses proteomics data to perform correlation network analysis and annotates protein-protein interactions, subcellular localizations and drug associations.

Resource Type: data processing software, software application, data analysis software, software resource

Keywords: Proteomics data, correlation network analysis, proteome wide systems genetics, genetic variants analysis, protein networks, phenotypic traits, protein-protein interactions, subcellular localizations, drug associations

Availability: Free, Freely available

Resource Name: CoffeeProt

Resource ID: SCR_022256

Record Creation Time: 20220506T050143+0000

Record Last Update: 20260815T182718+0000

Ratings and Alerts

No rating or validation information has been found for CoffeeProt.

No alerts have been found for CoffeeProt.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Molendijk J, et al. (2022) Proteome-wide systems genetics identifies UFMylation as a regulator of skeletal muscle function. eLife, 11.

Molendijk J, et al. (2021) CoffeeProt: an online tool for correlation and functional enrichment of systems genetics data. Nucleic acids research, 49(W1), W104.