

Resource Summary Report

Generated by [SPARC SAWG](#) on Jul 28, 2026

PHYTOPROT

RRID:SCR_013478

Type: Tool

Proper Citation

PHYTOPROT (RRID:SCR_013478)

Resource Information

URL: <http://urgi.versailles.inra.fr/phytoprot/>

Proper Citation: PHYTOPROT (RRID:SCR_013478)

Description: A database of protein domains shared by different sequence clusters. These were assembled by performing an "all-by-all" comparison of (putative) protein sequences with the BIOFACET software, building clusters of sequences based on their similarity and finally displaying ?? la Prodom the domains shared by the sequences in each cluster.

Synonyms: PHYTOPROT

Resource Type: database, data or information resource

Resource Name: PHYTOPROT

Resource ID: SCR_013478

Alternate IDs: nif-0000-03285

Record Creation Time: 20220129T080316+0000

Record Last Update: 20260725T191207+0000

Ratings and Alerts

No rating or validation information has been found for PHYTOPROT.

No alerts have been found for PHYTOPROT.

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 [SPARC SAWG](#) .

Coulibaly I, et al. (2008) Bioinformatic tools for inferring functional information from plant microarray data II: Analysis beyond single gene. International journal of plant genomics, 2008, 893941.